Ltd 5.1.8 Biocceleration version - 2006 using sw model GenCore (c) 1993 protein search, Copyright 1 protein Run

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updates/sec Search time 158.667 Seconds (without alignments) 391.900 Million cell updar

US-09-889-936A-6 737 score: Title: Perfect

.. WDYGTTYGYFDVWGQGTTVT MRVLILLWLFTAFPGILSDV Sequence:

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Scoring table

0.5 Gapext BLOSUM62 Gapop 10.0 residues 457216429 2589679 segs, Searched:

2589679 satisfying chosen parameters hits of number Total

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Post-processing:

Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database

Geneseq\_8:\*
geneseqp1980s:\*
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geneseqp2000s:\*
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geneseqp2003as:\*
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No. is the number of results predicted by chance to have a greater than or equal to the score of the result being printed, a derived by analysis of the total score distribution. score gand is Pred.

# SUMMARIES

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### ALIGNMENTS

5B3 antibody; IgG1; piezoelectric immunoassay; molecule explosive detection; 2,4,6-trinitrotoluene; region. chain variable standard; protein; 136 AA entry) antibody heavy (first 08-JAN-2001 **5B3** AAB15672 **AAB15672** Mouse; small n Murine gg RESULT 1 AAB15672 Mus 

WO200043774-A2

27-JUL-2000.

2000WO-IL000048 25-JAN-2000;

& DEV CO.
DEV CO LTD. 99IL-00128212 RES YISSUM RE YEDA RES 25-JAN-1999; (YISS ) (YEDA )

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2; Eshhar 'n Willner

WPI; 2000-524259/47 N-PSDB; AAA74604.

especially explosives comprises Apparatus for detecting small molecules, a piezoelectric crystal.

Disclosure, Fig 3A; 90pp; English

The present sequence is the heavy chain variable region of 583 antibody. The cDNA encoding this sequence was obtained from total RNA extracted from a 583 hybridoma by RT-PCR. 583 is an IgG1 antibody derived from a TNP-KLH immunised mouse. Its binding to TNP or DNS antigen can be blocked by very low amounts of TNT and it can therefore be used in a method for detecting small assayed explosive molecules. Molecules are detected using a piezoelectric sensor. Piezoelectric immunoassaying in liquid phase allows stationary and flow analysis of an aqueous sample. The method is sufficiently sensitive for detection of low molecular weight molecules

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which specifically binds to KID3, useful for treating
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call; the ability to bind to a portion of KiD3 that is exposed on the attacent of a living call in witro or in vivo; the ability to deliver a therapeutic agent or detectable marker to a cancer call expressing KiD3. Also included are an isolated mucleic acid sequence coding for the immunoglobulin polypeptide for its epitope muclei acid sequence coding for the immunoglobulin polypeptide or its epitope binding fragment, comprising with a vector containing the mucleic acid, producing a substantially purified immunoglobulin polypeptide or epitope binding fragment, comprising growing a cell line transferced/transformed/infected with a vector containing the mucleic acid, under conditions in which the immunoglobulin polypeptide or epitope binding fragment comprisising growing a cell line transferced/transformed/infected with the mucleic acid under conditions in which the immunoglobulin polypeptide or epitope binding fragment in golypeptide or epitope binding fragment, a pharmaceutical composition (comprising a therapeutically effective dese of the purified immunoglobulin polypeptide or tis epitope binding fragment, a polypeptide or the appropriate or tis epitope binding fragment, and a carrier), a pharmaceutical composition of comprising a therapeutically effective dose of the purified immunoglobulin or its epitope-binding fragment that specifically binds to kID3, and a context of an isolated call line consisting of American Type Culture or comprising the presence or absence of a cancer cells in an individual conference or absence or a pharmaceutical composition comprising a therapeutically effective dose of the agent, and a KID3 modulator. The conference or all single dabove is elected from an eracinoma, associated cancer (adamention and poly tumors, elliptions) bone cancer (adamentions, poly tumors, beland under cancer (adamentions, composition comprised and an individual carcinoma, colon cancer (adamentions, poly tumors, periforms, desemplastic fateses of the book cancer (adamentions, periforms, desemplastic manifor
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ADZ81889

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standard; protein;

entry) (first 14-JUL-2005 ΩÏ chain SEQ gamma2 heavy Anti-laminin-5

diagnosis; engineering; cytostatic; neoplasm; antibody chain. cancer; heavy antibody; Laminin;

Synthetic

WO2005040219

06-MAY-2005

2004WO-DK000744 28-OCT-2004

2003US-00695559. 2003WO-EP012012. 2003US-0523895P. 2004US-0567096P. 2004US-0568655P. 28-OCT-2003; 29-OCT-2003; 20-NOV-2003; 29-APR-2004; 06-MAY-2004; 13-MAY-2004;

AS BIOSTRATUM INC NORDISK NOVO (NOVO)

Salo တ Tarabykina SB, Padkar Mathiasen IS, Tryggvason K, Boutard A;

2005-355824/36 WPI;

b 5D5, mAb diagnosing, for competes with 6C12, useful that s and peptide t both 5D5 cancer nin-5 gamma2-binding cross-competes with or preventing laminin-5 treating,

English 338pp; 96; ID NO Disclosure;

The invention relates to a peptide that competes with mab 5D5, mab 6C12, or cross-competes with both 5D5 and 6C12 in specifically binding to at least a portion of domain III of the gamma2 chain of human laminin-5 (Ln-5 gamma2 DIII). Also included are a composition comprising an amount of the isolated peptide and a pharmaceutical carrier, an assay for assessing the gamma2-associated peptide content of a composition (comprising administering the peptide to the composition and determining whether the peptide binds any gamma2-associated peptides in the composition), a method of reducing cancer progression in a patient and a method of treating cancer in a patient. The peptide is a monoclonal antibody, preferably a humanized antibody or a fully human antibody. It binds a protein that is secreted from human cancer cells and comprises at least a portion of Ln-5 gamma2 DIII. The peptide is also conjugated to a compound the nucleic acid, and the composition are useful in the preparation of a medicament for the treatment of cancer. The peptides, nucleic acids, and the composition are useful for diagnosing, treating, or preventing cancer. The present sequence represents a Anti-laminin-5 gamma2 antibody heavy chain. t least a compound

The invention relates to an antibody or its fragment, capable of recognizing and binding to a three-dimensional structure of an extracellular region of a polypeptide encoded by a p53 apoptosis effector related to PMP-22 (PERP) gene. The invention also relates to a hybridoma that produces the antibody, a DNA encoding the antibody, a recombinant vector comprising the DNA, a host cell obtained by transducing the vector to the host cell, a method of producing the antibody and a therapeutic agent comprising the antibody as an active ingredient. The antibody is useful in an immunoassay for measuring the polypeptide encoded by the PERP gene, in the cell. The immunoassay is immunoprecipitation. The antibody is useful for measuring the expression of a polypeptide encoded by the PERP gene, using a fluorescent cell staining technique. The antibody is useful as a reagent for detecting the polypeptide encoded by the PERP gene and for diagnosing the disease associated with a

Novel antibody capable of binding to three-dimensional structure of extracellular region of polypeptide encoded by p53 apoptosis effector related to PMP-22 gene, useful for manufacturing diagnostic and therapeutic agent for cancer.

Japanese

SEQ ID NO 12; 121pp;

Claim 13;

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                                                    GNKLEWMGYISYSGFTSYNPSLRSRISFTRDTSKNQFFLQLNSVTSEDTATYYCARWDYG
                               to PMP-22
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                                                                                                                                                            sis; p53 apoptosis effect cytostatic; immunotherapy
      Score 622.5; DB 9;
Pred. No. 2.7e-50;
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a E, Sasaki
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                                                                                                                                                 related protein
                                                                                                                 standard; protein;
                                                                                                                                                            Antibody therapy; diagnosis
PERP; cancer; neoplasm; cyto
          84.5%
86.0%
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                                                                                                                                       entry)
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---NFFDYWGQGTTLT
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Ohta S, H
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JAPAN NAT C
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136
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polypeptide encoded by the PERP gene. The disease is cancer. The antibous useful in diagnosing a disease by detecting or measuring the amount a polypeptide encoded by the PERP gene in a cell, for treating cancer, for manufacturing a diagnostic, for detecting cancer and for manufacturing a therapeutic agent for cancer. This sequence represents murine PERP related protein of the invention. Note: The sequence data this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
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49. .54
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ilarity 81.6%;
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of myocard
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                                                                                                                                   Local Sim
hes 111;
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Claim 6; Fig

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         Monoclonal antibodies which inhibit type II phospholipase A2 are useful in the treatment of myocardial infarction, cerebral infarction, acute kidney failure, chronic rheumatism, cardiac shock, pancreatitis, adult respiratory distress syndrome and colitis. The antibodies were generated by immunising Balb/C mice with recombinant human type II phospholipase A2. Spleen cells from the mice were fused with mouse myeloma P3U1 (P3x63Ag8.U1) and the hybridomas obtained were screened for phospholipase A2 inhibitory activity. Active clones were isolated including 12H5, 1.4 and 10.1. These were cultured and the antibody isolated from the culture supernatant by precipitation with ammonium sulphate and purification on a column of protein A-Sepharose CL4B. Because the antibody acts on the primate and mouse forms of enzyme as well as human it is particularly suitable for preclinical testing
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                                                                                                                                                                                                                                                                                                              cisplatin;
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II phospholipase A2 activ
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                                                                                                                                                                                                                                     11;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     type II phospholipase A2; inhibition; disorder; nephrotoxicity; anticancer.
                                                                                                                                                                                                             Length
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                                                                                                                                                                                                      Score 588.5; DB 2;
Pred. No. 4.1e-47;
. Mismatches 11;
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oody inhibiting
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                                                                                                                                                                                                            79.9%;
ilarity 81.4%;
Conservative
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96JP-00247635
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with an antibody
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on; kidney
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      standard;
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114; Conser
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N-PSDB; AAV12261
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19-SEP-1996;
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disorders (such as acute renal failure) associated with the
administration of cisplatin for the treatment of cancer. The method
comprises treatment with a monoclonal antibody which inhibits the
activity of type II phospholipase A2 (particularly of type II
phospholipase A2 of human origin), or with a protein or peptide
possessing the same inhibitory activity and containing a part of the
antibody sequence. Preferably the antibody also inhibits the activity of
antibody sequence. Preferably the antibody also inhibits the activity of
cype II phospholipase A2 bound to a cell membrane. Three specific
cype II phospholipase A2 bound to a cell membrane. Three specific
monoclonal antibodies having these properties which can be used are 12H5,
C 10.1 and 1.4, derived from hybridomas FERM BP-5300, FERM BP-5298 and FERM
C BP-5297 respectively. The method can be used for suppressing the
c nephrotoxicity which is a characteristic feature of cisplatin
c administration, and therefore allowing more efficient use of this drug as
c an anticancer agent, e.g. by allowing an increased dosage to be used
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heavy chain; variable region; mouse; monoclonal; hybridoma; PM1; plasmid
pPM-k3; pPM-h1.
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Pred. No. 4.1e
4; Mismatches
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92JP-00032084
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11 Similarity 81.4%;
114; Conservative
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(first entry)
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19. .137
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N-PSDB; AAQ30756.
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9-FEB-1992;
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30-MAR-1993
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                                                  The sequences given in AAR28670-71 were encoded by plasmids which were used in example to illustrate the production of a human antibody which recognises human interleukin-6 receptor (IL-6R). The antibody comprises light (L) chain and heavy (H) chain variable regions which were derived from a mouse monoclonal antibody produced from the hybridoma PM1 which contained the plasmids pPM-k3 and pPM-h1. (Updated on 25-MAR-2003 to
        interleukin-6 receptor - has low
on complementarily determining
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Cavenee WK,
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2001US-0342258P.
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Nice EC
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28-SEP-2001;
21-DEC-2001;
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                                        Disclosure;
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The invention relates to an isolated specific binding member (I) which recognizes an epidermal growth factor receptor (EGFR) epitope which is found in tumourigenic, hyperproliferative or abnormal cells and not detectable in normal cells. The EGFR epitope is located within the region comprising residues 273-501 of EGFR and does not demonstrate any amino acid sequence alterations or substitutions from normal EGFR. (I) is capable of binding the de2-7 EGFR and absence of amplification of junctional peptide and does not bind to EGFR on normal EGFR. (I) is absence of aberrant expression, or in the absence of amplification of normal wild-type gene. The binding member comprises the VH and VL polypeptide sequence of monoclonal antibody (mAb) 806. (I) is useful for treating or diagnosing human or animal body, especially for treating treating or diagnosing human or animal body, especially for treating treating or treating cancer which is located in or adjacent the brain, in a mammal. (I) is useful for the preparation of amedicament and for preventing or treating cancer which is located in or adjacent the brain, in a mammal. (I) is also useful for detecting the presence of amplified EGFR, de2-7EGFR or EGFR with high mannose glycosylation, where the presence of amplified EGFR, de2-7EGFR or EGFR with a high mannose glycosylation of suspicing a biological sample from a mammal in which the presence of amplified EGFR from the sample and the antibody, where the detecting the presence or activity of an EGFR which indicates the expression of binding indicates that presence or activity of the EGFR will be amplified EGFR from the mammal. A pharmaceutical expression or concer in the mammal. A pharmaceutical expression or concer in the mammal and optionally vehicle, carrier or dilluent is composition composition or proper in the mammal of account or definite and optionally vehicles.
                                                              cells
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             particularly antibodies recognizing repitope found only in tumorigenic ag and treating cancer in mammals.
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and/or treating cancer
  specific binding members, panal growth factor receptor efor diagnosing, preventing
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Novel specific kepidermal growthuseful for diagr
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GNKLEWMGYISYSGFTSYNPSLRSRISFTRDTSKNQFFLQLNSVTSEDTATYYCARWDYG MRVLILLWLFTAFPGILSDVQLQESGPGLVKPSQSLSLTCSVTGYSITSGYAWNWIRQFP Length 134; Indels i7; 14; .. 6 2e-47 B ed. No. 6.20 Mismatches Score 586.5; Pred. No. 6. 4. 136 131 79.6%; ilarity 83.1%; Conservative ----FPYWGQGTLVT TTYGYFDVWGQGTTVT l Similarity 113; Conser Ž Н Match 61 121 121 Local Best Loc Matches Query <u>გ</u> 20 ઠે g õ δ

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the invention standard; protein; to entry) protein relating (first 16-DEC-2004 ADS14282; Murine ADS14282 RESULT 

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endocrine; thrombocytopenia receptor; antidiabetic; haemostatic; endo diabetes; laron syndrome; thrombopoietin. variant type II

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antidiabetic; haemostatic; endocrine; laron syndrome; thrombopoietin.

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                                                                                                                                                                                                                                                                                              having agonist colored ligand (preferably an antibody) (I) having agonist activity with respect to a variant receptor. A ligand of the invention has antidiabetic, haemostatic, and endocrine-gen. activity. The ligand is useful for treating the disease caused due to variant receptor, which involves combining the ligand/antibody with the variant receptor, where the disease is congenital non-megakaryocyte thrombocytopenia. A ligand of the invention is useful for transmitting signal to a variant receptor, which involves combining the variant receptor with the ligand. The invention is useful for treating diseases resulting due to variation receptor, such as type II diabetes and laron syndrome. The ligand enables signal transduction function of thrombopoietin, with respect to variant receptor. The present sequence i used in the exemplification of the invention.
                                                                                                                                                                                                          Novel ligand having agonist activity with respect to variant receptor e.g., variant thrombopoietin receptor, useful for treating disease resulting due to variant receptor such as congenital non-megakaryocyte
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Pred. No. 5e-45
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N-PSDB; ADS14281.
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108; Conserv
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                                                                                                                                                    Novel ligand having agonist activity with respect to variant receptor e.g., variant thrombopoietin receptor, useful for treating disease resulting due to variant receptor such as congenital non-megakaryocyte
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disease;
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N-PSDB; ADS14287.
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3-SEP-2004.
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                                                                                                                                                                                     which specification describes a monoclonal antibody, designated anti-C3-2, which specifically binds on a functional domain exposed on native human C3, thus inhibiting the generation of the biologically active peptides such as C3a and C5a. The functional domain is in part located on the 23 kD-a-chain fragment of C3c. The antibody is useful for the preparation of a medicament for inhibiting complement activation, preferably for preventing or treating inflammatory diseases mediated by activation of complement. The present sequence represents the consensus sequence of the heavy chain variable region of the monoclonal antibody anti-C3-2, derive from N-terminal sequencing, primer-induced sequence and sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        The specification describes a monoclonal antibody, designated anti-C3-2, which specifically binds on a functional domain exposed on native human C3, thus inhibiting the generation of the biologically active peptides such as C3a and C5a. The functional domain is in part located on the 23 kD-a-chain fragment of C3c. The antibody is useful for the preparation of a medicament for inhibiting complement activation, preferably for preventing or treating inflammatory diseases mediated by activation of complement. The present sequence represents the heavy chain region of the monoclonal antibody anti-C3-2.
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Ind diagnosis
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Pred. No. 3.5e
3; Mismatches
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                                                                                                                                                                New molecule capable of inhibiting specifically binds on a functional useful for preventing or treating i
                                                                                                                                                                                                                                                                                                                                                                                    Disclosure, Fig 14B; 72pp; English
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useful for treatment, prevention
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fragments, that bind to human insulin-like growth factor-1 receptor (IGF-IR) and optionally: (i) inhibit natural binding of insulin-like growth factors (IGF)-1 and/or -2; and/or (ii) inhibit specifically tyrosine factors (IGF)-1 and/or -2; and/or (ii) inhibit specifically tyrosine chinase activity of IGF-IR. Ab and its fragments are used to prevent or treat diseases associated with overexpression and/or abnormal activity of IGF-IR and/or epidermal growth factor receptor (EGFR) and/or with chese receptors with their ligands. Especially they inhibit transformation of normal cells to tumor cells, inhibit growth and/or proliferation of tumor cells, so are useful against cancers of the prostate, lung, breast, endometrium and colon, also osteosarcoma, and also for treating psoriasis. Ab are also used to diagnose diseases caused by abnormal expression of IGF-IR and/or EGFR. This sequence represents a protein sequence used to generate the Ab of the invention.
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The invention relates to a novel isolated anti-insulin-like growth factor capable of pinding to human IGF-IR and, if necessary, capable of capable of binding to human IGF-IR and, if necessary, capable of expectically inhibiting tyrosine kinase activity of the receptor, comprising a light or heavy chain having at least one complementary determining region (CDR) consisting of one of two fully defined 16 amino acids (ADSG-7006 and ADSG-7014). An antibody of the invention is useful in the preparation of a medicament intended for the prevention or treatment of an illness connected with an overexpression and/or an abnormal appearation of the IGF-IR and/or connected with a hyperactivation of the transduction pathway of the signal mediated by the induces secondary effects connected with inhibition of the insulin receptor. The antibody is useful for preparation of a medicament intended to inhibit the transformation of normal cells into cells with tumoral character, preferably IGF-dependent, especially IGF1 and/or IGF2. dependent and/or HERZ/neu-dependent and/or EGF-dependent and/or HERZ/neu-dependent cells. (I) is useful for preparation of a medicament intended to inhibit the growth and/or the projected prevention of tumor cells, preferably IGF-dependent and/or IGF2 dependent and/or IGF2-dependent and/or IGF2-dependent and/or IGF2-dependent and/or IGF2-dependent and/or IGF2-dependent and/or IGF2-dependent cells. (I) is useful in the preparation of a medicament intended for the prevention or for the treatment of psoriasis. (I) is useful in the preparation of a medicament intended for the prevention of a medicament intended for the specific targeting of encorer, osteosarcoma, lump cancer, the cancer, endometrial cancer or colon cancer. (I) is useful in the preparation of a medicament intended for the specific targeting the prevention of a medicament intended for the specific targeting the prevention of a medicament intended for the specific targeting the prevention or for the treatment of psoriasis of intended for the pre
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                                                                                                                                                                                                                                                                                                                     novel isolated anti-insulin-like growth factor I receptor (IGF-IR) antibody or its functional fragment, being capable of binding human IR and specifically inhibiting tyrosine kinase activity of receptor, useful for treating cancer.
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The invention relates to a novel humanized anti-insulin like growth factor I receptor (IGF-IR) antibody (Ia), anti-insulin/IGF-5 hybrid receptor (hybrid-R) antibody (Ib), or anti-IGF-IR and hybrid-R antibody (Ic) or their antigen-binding fragments, comprising a complementary determining region of non-human origin and a framework region having a human residue, and capable of inhibiting in vivo tumor growth. A composition of the invention has cytostatic activity. An antibody of the invention is useful for determining regression progression or onset of a activation of human IGF-IR, hybrid-R or both IGF-IR and hybrid-R relative to normal. A pharmaceutical composition of the invention is useful for the treatment or prevention of a disorder chosen from breast cancer, chosen cancer, kidney cancer, thyroid cancer, osteosarcoma, panciestic cancer, and myeloma. The cancer, kidney and myeloma. The n isolated from region lung cancer, bla a, pancreatic can chain variable equence represents a heavy 7010 cancer, thyroid cancer, present sequence antibody monoclonal

receptor ce, of IGF-

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A. 127 Sequence

Query Match

Length φ. .. OB 54; ហ Score 5.2%;

Search completed: May 19, 2006, 17:40:02 Job time : 161.667 secs

5.1.8 Biocceleration GenCore version (c) 1993 - 2006 Copyright

using sw model protein search, protein Search time 25.9048 Seconds (without alignments) 505.138 Million cell upday 17:40:37 2006, May 19,

Run on:

updates/sec

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US-09-889-936A-6 737

.... WDYGTTYGYFDVWGQGTTVT MRVLILLWLFTAFPGILSDV score: Title: Perfect so Sequence:

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residues 96216763 283416 segs, Searched:

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

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18-Oct-1

RESULT 2
B24672
Ig heavy chain precursor V region (VGAM3-2) - mouse
C; Species: Mus musculus (house mouse)
C; Date: 19-Nov-1988 #sequence\_revision 19-Nov-1988 #text\_change
C; Accession: B24672
C; Accession: B24672
R; Winter, E.; Radbruch, A.; Krawinkel, U.
EMBO J. 4, 2861-2867, 1985
A; Reference number: A91022; MUID: 86055722; PMID: 2998759

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Best Local Similarity 82.4%; Pred. No. 2.6e-45
Matches 112; Conservative 12; Mismatches
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C; Date: 24-Apr-1984 #sequence revision 30-Jun-1992 #text change C; Accession: PL0102; S03262; A93814; A91462; A93787; S23599
R; Rinfret, A.; Horne, C.; Dorrington, K.J.; Klein, M.
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A; Accession: B24672
A; Molecule type: DNA
A; Residues: 1-134 <WIN>
A; Cross-references: UNIPARC:UPI00001768F5
A; Note: this sequence was determined from the differentiat C; Genetics:
A; Introns: 15/1
C; Superfamily: immunoglobulin V region; immunoglobulin hom C; Keywords: heterotetramer; immunoglobulin
F; 19-134/Product: Ig heavy chain V region VGAM3-2 #status F; 33-116/Domain: immunoglobulin homology <IMM>
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C; Species: Mus musculus (house mouse)
C; Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text_c
C; Accession: S30752
R; Grant, F.J.; Levin, S.D.; Gilbert, T.; Kindsvogel, W.
Nucleic Acids Res. 15, 5496, 1987
A; Title: Improved RNA sequencing method to determine immun
A; Reference number: S30751; MUID:87260030; PMID:3601683
A; Accession: S30752
A; Molecule type: mRNA
A; Residues: 1-149 <GRA>
A; Cross-references: UNIPARC:UPI0000115D92; EMBL:X05878; NI
C; Superfamily: immunoglobulin V region; immunoglobulin hom
C; Keywords: heterotetramer; immunoglobulin
F; 33-116/Domain: C region (C-gamma 2b) (fragment) #status
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Pred. No. 5.4e-43;
5; Mismatches 15
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A, Richert cloning, sequencing and expression of the rearranged MOPC 315 VH gene A, Reference number: PLUID2; WILL:9923951; PMID:2497341
A, Arcession: PLUID2; MILL:9923951; PMID:2497341
A, Arcession: PLUID2; MILL:9923951; PMID:2497341
A, Restance tryes: ANN PROPERTY, June 1988
A, Regerence number: S01262
A, Molecule type: DNA
A, Restance number: S01262
A, Molecule type: DNA
A, Restance number: S01262
A, Molecule type: DNA
A, Restance number: A512, G'. Le 137
A, RESTANCE A, DNA
A, Restance number: A514, M. A, S692-5696, 1977
A, Restance number: A514, M. A, S692-5696
A, Restance number: A5
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ced. No. 1.8e-41;
Mismatches 16
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C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 20-Jun-2
C;Accession: S38718
R;Cimanis, A.Y.
submitted to the EMBL Data Library, November 1993
A;Reference number: S38713
A;Accession: S38718
A;Accession: S38718
A;Accession: S38718
A;Molecule type: mRNA
A;Kesidues: 1-116 <CIM>
A;Residues: 1-116 <CIM>
A;Cross-references: UNIPARC:UPI0000117542; EMBL:X76018; NID:g416102; PID
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology <IMM>
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Ig heavy chain precursor V region (M315) - mouse
C; Species: Mus musculus (house mouse)
C; Date: 30-Jun-1990 #sequence_revision 30-Jun-1990 #text_change 09-C; Accession: JT0509
C; Accession: JT0509
R; Levy, N.S.; Malipiero, U.V.; Lebecque, S.G.; Gearhart, P.J.
J. Exp. Med. 169, 2007-2019, 1989
A; Title: Early onset of somatic mutation in immunoglobulin VH genes
A; Reference number: JT0501; MUID: 89279149; PMID: 2499654
A; Accession: JT0509
A; Ratus: translation not shown
A; Molecule type: mRNA
A; Residues: 1-116 < LEV>
A; Cross-references: UNIPROT: P18531; UNIPARC: UP100000278E0
A; Experimental source: strain BALB/cJ
C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
F;1-18/Domain: signal sequence #status predicted
F;19-116/Pomain: immunoglobulin homology <IMM>
F;33-116/Domain: immunoglobulin homology <IMM>
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73.1%; Score 539; DB 1; Best Local Similarity 87.1%; Pred. No. 1.2e-39; Matches 101; Conservative 7; Mismatches 8;
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Residues: 1-116 <LEV>
Cross-references: UNIPROT:P18532; UNIPARC:UPI00000278E1
Cross-references: UNIPROT:P18532; UNIPARC:UPI00000278E1
Experimental source: strain BALB/cJ
Note: this sequence belongs to the VH3660 subfamily
Superfamily: immunoglobulin V region; immunoglobulin
Keywords: heterotetramer; immunoglobulin
1-18/Domain: signal sequence #status predicted <SIG>
1-18/Domain: immunoglobulin homology <IMM>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ig heavy chain V region (PTF.02) - mouse C; Species: Mus musculus (house mouse) C; Date: 07-Sep-1990 #sequence_revision 07-Sep-1990 #text_char C; Accession: S07637  
R; Urakov, D.N.; Deev, S.M.; Polyanovsky, O.L. Nucleic Acids Res. 17, 9481, 1989  
A; Title: The structure of the expressible VH gene from a hybra A; Reference number: S07637; MUID: 90067954; PMID: 2587273  
A; Molecule type: DNA A; Residues: 1-136 <URA. A; Residues: 15/3  
C; Genetics: A; Introns: 15/3  
C; Superfamily: immunoglobuli- Tark for residue 112  
C; Keywords: heterre
                                                                                                                                                                                                                                          immunoglobulin VH
PMID:2499654
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Pred. No. 3.3e-41;
5; Mismatches 5;
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                                                                                                                                 mouse
                                                                                                                                                        C; Species: Mus. musculus (nouse mouse)
C; Date: 30-Jun-1990 #sequence_revision 30-Jun
C; Accession: JT0508
R; Levy, N.S.; Malipiero, U.V.; Lebecque, S.G.
J. Exp. Med. 169, 2007-2019, 1989
A; Title: Early onset of somatic mutation in in
A; Reference number: JT0501; MUID:89279149; PM
A; Reference number: JT0501; MUID:89279149; PM
A; Reference number: JT0508
A; Status: translation not shown
A; Molecule type: mRNA
A; Molecule type: mRNA
A; Residues: 1-116 < LEV>
A; Cross-references: UNIPROT:P18532; UNIPARC:U
A; Experimental source: strain BALB/cJ
A; Note: this sequence belongs to the VH3660 s'
C; Superfamily: immunoglobulin V region; immunoC; Keywords: heterotetramer; immunoglobulin
C; Keywords: heterotetramer; immunoglobulin
F; 1-18/Domain: signal sequence #status predic
F; 19-116/Product: Ig heavy chain V region (1BF)
F; 33-116/Domain: immunoglobulin homology < IMM
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(house mouse)
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ilarity 91.4%;
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Schilarity 77.9%;
5, Conservative
TTYGYFDVWGQGTTVT
            Ig heavy chain precursor V reg
C;Species: Mus musculus (house
C;Date: 30-Jun-1990 #sequence_
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Cross-references: UNIPARC:UPI0000115D24; GB:X03378; NID:g52007; PIDN:CAA27095.1;
Superfamily: immunoglobulin V region; immunoglobulin homology
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Mol. Immunol. 28, 1063-1072, 1991
A; Title: Molecular characterization of monoclonal anti-steroid antibodies and their pH-reactivity profiles.
A; Reference number: A53285; MUID:92017897; PMID:1922102
A; Accession: C53285
A; Status: preliminary
A; Molecule type: mRNA; protein
A; Residues: 1-119 <SAW>
A; Residues: 1-119 <SAW>
A; Cross-references: UNIPARC:UPI000011D0A5; GB:D12734; NID:9220548; PIDN:EA; Note: sequence extracted from NCBI backbone (NCBIN:63297, NCBIP:63302)
C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
F; 15-98/Domain: immunoglobulin homology <IMM>
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Cross-references: UNIPARC:UPI0000117638; EMBL:S54194; NID:g264864
Superfamily: immunoglobulin V region; immunoglobulin homology
15-98/Domain: immunoglobulin homology <IMM>
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T01262
Ig heavy chain V region - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 12-Feb-1999 #sequence_revision 12-Feb-1999 #text_cl
C;Accession: T01262
R;Pirofski, L.A.; Thomas, E.K.; Scharff, M.D.
AIDS Res. Hum. Retroviruses 9, 41-49, 1993
A;Title: Variable region gene utilization and mutation in 8
A;Reference number: Z14285; MUID:93152285; PMID:7678971
A;Reference number: Z14285; MUID:93152285; PMID:7678971
A;Residues: 1-114 <PIR>
A;Residues: 1-114 <PIR>
A;Residues: 1-114 <PIR>
A;Cross-references: UNIPARC:UPI0000117638; EMBL:S54194; NISCOSS-references: UNIPARC:UPIO000117638; EMBL:S54194; NISCOSS-references: UNIPARC:UPIO000117638; EMBL:S54194; NISCOSS-FEMENTALINE Immunoglobulin V region; immunoglobulin home
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E25114
Ig heavy chain V region (HP25) - mouse
C; Species: Mus musculus (house mouse)
C; Date: 29-Aug-1987 #sequence_revision 29-Aug-1987 #text_C; Accession: E25114
R;Ollier, P.; Rocca-Serra, J.; Somme, G.; Theze, J.; Fouse EMBO J. 4, 3681-3688, 1985
A; Title: The idiotypic network and the internal image: postities number: A91028; MUID:86136012; PMID:3937730
A; Accession: E25114
A; Molecule type: mRNA
A; Residues: 1-119 <OLL>
A; Cross-references: UNIPARC:UP10000115D24; GB:X03378; NIIC C; Superfamily: immunoglobulin but the immunoglobulin house.
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Pred. No. 4.6e-37
; Mismatches 1:
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Pred. No. 4.8e-37
5; Mismatches
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ilarity 82.2%;
Conservative
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Les: Mus musculus (house mouse)

102-May-1994 #sequence_revision 18-Nov-1994 #text_change

102-May-1994 #sequence_revision 18-Nov-1994 #text_change

103-May-1994 #sequence_revision 18-Nov-1994 #text_change
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Superfamily: immunoglobulin V region; immunoglobulin homology
15-98/Domain: immunoglobulin homology <IMM>
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 S37200

Ig heavy chain V region - mouse
C; Species: Mus musculus (house mouse)
C; Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_C; Accession: S37200
R; Fischer, R.; Voss, A.; Hunziker, W.; Stierhof, Y.D.; Kr submitted to the EMBL Data Library, August 1993
A; Description: Production and cloning of TMV-specific mon A; Reference number: S37200
A; Reference number: S37200
A; Accession: S37200
A; Status: preliminary
A; Molecule type: mRNA
A; Residues: 1-121 <FIS>
A; Cross-references: UNIPARC: UPI00001161AC; EMBL: X74587; N C; Superfamily: immunoglobulin V region; immunoglobulin horizoners immunoglobulin homology <IMM>
                                                              #text
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PMID:3267217
                                                                                               Y.D.;
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Pred. No. 3.4e-38;
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ed. No. 2.4e-38;
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C; Date: 01-Dec-1989 #sequence_revision 30-Sep
C; Accession: 128195
R; Sherman, M.A.; Deans, R.J.; Bolger, M.B.
J. Biol. Chem. 263, 4059-4063, 1988
A; Title: Haloperidol binding to monoclonal an A; Reference number: A28195; MUID:88153717; PM A; Accession: 128195
A; Molecule type: mRNA
A; Residues: 1-117 < SHE>
A; Cross-references: UNIPARC:UPI0000114D72; GB A; Note: the authors translated the codon AAC C; Superfamily: immunoglobulin V region; immun C; Keywords: heterotetramer; immunoglobulin F; 15-98/Domain: immunoglobulin homology < IMM>
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C; Species: Mus musculus (house mouse)
C; Date: 29-Aug-1987 #sequence_revision 29-Aug-1987 #text_change 20-Jun-2000
C; Date: 29-Aug-1987 #sequence_revision 29-Aug-1987 #text_change 20-Jun-2000
C; Accession: F2514
R; Ollier, P :, Rocca-Serra, J :, Somme, G :; Theze, J :; Fougereau, M.
EMBO J . 4, 3681-3688, 1985
A; Title: The idiotypic network and the internal image: possible regulation of a A; Reference number: A91028; MUID:86136012; PMID:3937730
A; Accession: F25114
A; Molecule type: mRNA
A; Residues: 1-115 - OLL.
A; Cross-references: UNIPARC:UPI0000115cn-
C; Superfamily: immunoglobulic.
C; Keywords: heterre.
F; 15-co.'-
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C;Species: Mus musculus (house mouse)
C;Date: 09-Mar-1990 #sequence_revision 09-Mar-1990 #text_change 23-C;Accession: D33932
C;Accession: D33932
R;Baccala, R.; Vo Quang, T.; Gilbert, M.; Ternynck, T.; Avrameas, S Proc. Natl. Acad. Sci. U.S.A. 86, 4624-4628, 1989
A;Title: Two murine natural polyreactive autoantibodies are encoded A;Tetle: Two murine natural polyreactive autoantibodies are encoded A;Accession: D33932
A;Accession: D33932
A;Accession: D33932
A;Ketus: preliminary
A;Molecule type: mRNA
A;Residues: 1-115 cAAC>
A;Cross-references: UNIPARC:UPI00001768F1; GB:M27106
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: immunoglobulin
F;33-115/Domain: immunoglobulin homology <IMM>
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16;
                                               Score 504; DB 2; ]
Pred. No. 1.2e-36; 5; Mismatches 13;
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Pred. No. 8.6e-36;
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fords: heterotetramer; immunoglobulin
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> Gapext BLOSUM62 Gapop 10.0 Scoring table:

925015592 residues 2849598 segs, Searched:

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2849598 hits satisfying chosen parameters: of Total number

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Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

UniProt\_7.2:\*
1: uniprot\_sprot:\*
2: uniprot\_trembl:\* Database

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

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758; PubMed=428562;
Wolf O., Zemell R., Burstein
d function of immunoglobulin g:1839-1845(1979).
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MEDLINE=78094475; PubMed=414225;

Jilka R.L., Pestka S.;

"Amino acid sequence of the precursor reimmunoglobulin heavy chain.";

Proc. Natl. Acad. Sci. U.S.A. 74:5692-56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-MAR-2006, entry version 51.

Ig heavy chain V region MOPC 315 precus
Mus musculus (Mouse).

Eukaryota; Metazoa; Chordata; Craniata
Mammalia; Eutheria; Euarchontoglires; (Muroidea; Muridae; Murinae; Mus.
  Commons
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MEDLINE=74170779; PubMed=4524622;
Francis S.H., Leslie R.G.Q., Hood L.,
                          mRNA
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UNKNOWN
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                                               Ensembl; ENSMUSGO0000054328; Mu
InterPro; IPR003599; Ig.
InterPro; IPR003110; Ig-like.
InterPro; IPR003597; Ig_cl.
InterPro; IPR003006; Ig_MHC.
InterPro; IPR013106; V-set.
Pfam; PF07654; Cl-set; 2.
SMART; SM00409; IG; 2.
SMART; SM00409; IG; 2.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 4.
PROSITE; PS00290; IG_MHC; UNKNOSEQUENCE 479 AA; 51992 MW;
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                         EMBL; BC002091; AAH02091.1;
HSSP; P01820; 1G7J.
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les 108; Conservative
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MEDLINE=79148758; PubMed=Schechter I., Wolf O., Ze "Structure and function of Fed. Proc. 38:1839-1845(1)
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Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

A Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

A Nillalon D.K., Manny D.W., Green E.J., Lu X., Gibbs R.A.,

A Nillalon A., Young A.C., Shevchenko Y., Bouffard G.G.,

A Richards A.C., Grimwood J., Schmutz J., Myers R.M.,

Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

A Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,

A Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;

"Generation and initial analysis of more than 15,000 full-length human
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi
Muroidea; Muridae; Murinae; Mus.
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Pred. No. 2.4e-50;
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SMART; SM00409; IG; 2.
SMART; SM00407; IGc1; 2.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG LIKE; 4
PROSITE; PS00290; IG MHC; UN
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MEDLINE=89238351; PubMed=2497341; DOI=10.1016/0161-5890(89)90133-8
Rinfret A., Horne C., Dorrington K.J., Klein M.;
"Cloning, sequencing and expression of the rearranged MOPC 315 VH gegment.";
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   (alpha) chain
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Complementarity-determining-2.
Framework-3.
Complementarity-determining-3.
Framework-4.

By similarity.
G -> GG (in Ref. 1; CAA30727).
G -> YG (in Ref. 2).
GY -> YG (in Ref. 4).
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Framework-1.
Complementarity-determining-1
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2).
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"Amino-acid sequence of the variable region of the heavy of a mouse myeloma protein with anti-hapten activity."; Proc. Natl. Acad. Sci. U.S.A. 71:1123-1127(1974).
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                               MEDLINE=77244979; PubMed=268248;
MEDLINE=77244979; PubMed=268248;
Padlan E.A., Davies D.R., Pecht I., Givol D., Wright C "Model-building studies of antigen-binding sites: the site of mopc-315.";
Cold Spring Harb. Symp. Quant. Biol. 41:627-637(1977).
-!- MISCELLANEOUS: This alpha chain was isolated from protein that has anti-dinitrophenyl activity.
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                                                                                                                                                                            Ensembl; Ensembl; Ensembl; Ensembl; Ensembl; Ensembl; Ig.

InterPro; IPR003599; Ig.
InterPro; IPR003596; Ig.v.
InterPro; IPR013106; V-set.
Pfam; PF07686; V-set; 1.
SMART; SM00406; IG; 1.
SMART; SM00406; IG; 1.
PROSITE; PS50835; IG_LIKE; 1.
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EMBL; X07880; CAA30727.1; -; Genomic_DNA.
PIR; PL0102; AVMS35.
HSSP; P01820; 1G7J.
SMR; P01822; 20-137.
Ensembl; ENSMUSG0000057048; Mus musculus
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TISSUE=Spleen;

MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;

X Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

A Strausberg R.L., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schemen C.M., Schuler G.D.,

A Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

Raha S.S., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

Nillalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

Raha J., Helton B., Ketteman M., Madan A., Young A.C., Shevchenko Y., Boutfard G.G.,

Whiting M., Madan A., Young A.C., Shevchenko Y., Boutfard G.G.,

Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,

Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,

"Generation and initial analysis of more than 15,000 full-length human
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Distributed under the Creative Commons Attribution-NoDerivs License
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                                                                                                    Rattus norvegicus (Rat).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Muridae; Murinae; Rattus.
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NIH MGC Project;
Submitted (APR-2005) to the EMBL/GenBank/DDBJ databases
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BE5C2483C69F186C CRC64;
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id. Sci. U.S.A. 99:16899-16903(2002)
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Pred. No. 1.6e-45
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InterPro; IPR003599; Ig.
InterPro; IPR003110; Ig-like.
InterPro; IPR003597; Ig_c1.
InterPro; IPR003596; Ig_v.
InterPro; IPR013106; V-set.
Pfam; PF07654; C1-set; 4.
SMART; SM00409; IG; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 5.
PROSITE; PS00290; IG_MHC; UNKNOWN 3.
SEQUENCE 615 AA; 67986 MW; BE5C248
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10-MAY-2005, sequence version 07-FEB-2006, entry version 12. LOC314509 protein. Name=LOC314509;
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80.3%;
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5, sequence version 1.
6, entry version 13.
protein.
                                                                                                   into UniProtKB/TrEMBL
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                                                                                                                                                                                                                                                                                     Mus
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SMR; Q6LBQ5; 20-136.

Ensembl; ENSMUSG0000057048; Mt
InterPro; IPR003599; Ig.
InterPro; IPR003596; Ig.v.
InterPro; IPR013106; V-Set.
SMART; SM00409; IG; 1.
PROSITE; PS50835; IG_LIKE; 1.
Immunoglobulin domain.
NON_TER
                                                                                                         05-JUL-2004, sequence version 07-FEB-2006, entry version 11. VH gene product (Fragment).
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SMR; Q6LBQ5; 20-136
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                                                                                  PRELIMINARY;
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il Similarity 77.9%;
106; Conservative
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QS MOUSE
QGLBQS;
QGLBQS;
05-JUL-2004, 8
05-JUL-2004, 8
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Q569B8;
10-MAY-2005, i
10-MAY-2005, e
07-FEB-2006, e
RGD1359202 pro
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Н
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Distributed under the Creative Commons Attribution-NoDerivs License
                                                                           Euteleostomi;
ia; Sciurognathi;
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                                                                                                                                                                                                                                                                                            region;
                                                                                                                                 DOI=10.1084/jem.169.6.2007
que S.G., Gearhart P.J.;
in immunoglobulin VH genes
                                                                                                                                                                                                                                                                                                          Ig heavy chain V region 1B43.
/FTId=PRO_0000015242.
Framework-1.
Complementarity-determining-1
Framework-2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11
                                                                                                                                                                                                            PIR; JT0508; HVMS1B.

PDB; IKCS; X-ray; H=25-116.

PDB; IKCV; X-ray; H=25-116.

Ensembl; ENSMUSG0000061612; Mus musculus.

InterPro; IPR007110; Ig-like.

InterPro; IPR013106; V-set.

Pfam; PF07686; V-set; 1.

SMART; SM00406; IGv; 1.

PROSITE; PS50835; IG_LIKE; 1.

3D-structure; Immunoglobulin domain; Immunoglobulin V region 1B43

SIGNAL 19 116 Ig heavy chain V region 1B43
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                  HV61 MOUSE STANDARD; PRT; 116 AA.
P18532;
01-NOV-1990, integrated into UniProtKB/Swiss-Prot.
01-NOV-1990, sequence version 1.
07-MAR-2006, entry version 44.
Ig heavy chain V region 1843 precursor.
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eute Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
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Pred. No. 6.1e-46
; Mismatches
                                                                                                                       STRAIN=BALB/cJ;
MEDLINE=89279149; PubMed=2499654; DOI=10.108
Levy N.S., Malipiero U.V., Lebecque S.G., Ge
"Early onset of somatic mutation in immunogl
primary immune response.";
J. Exp. Med. 169:2007-2019(1989).
-!- MISCELLANEOUS: This sequence belongs to
                                                                                                                                                                                                                                                                                                                                                                  Bimilarity
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larity 91.4%;
Conservative
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                                                                                                  NCBI_TaxID=10090;
[1]
NUCLEOTIDE SEQUEN
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Shas musculus (Mouse).

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Chammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;

Charolea; Murinae; Mus.

NCBI_TaxID=10090;

NCCBI_TaxID=10090;

NCCBI_TaxID=10090
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116
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TISSUE=Spleen;

TISSUE=Spleen;

A Figuraberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

A Klausner R.D., Colling F.S., Wagner L., Shenmen C.M., Schuler G.D.,

A Lischul.S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

Brownstein M.J., Usdin T.B., Former A.A., Rubin G.M., Hong L.,

Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

Brownstein M.J., Wafley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

A Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

A Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,

Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,

A Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.,

"Generation and initial analysis of more than 15,000 full-length human and mouse cDNA sequences.";
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Distributed under the Creative Commons Attribution-NoDerivs License
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Elikaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Muridae; Murinae; Rattus.

NCBI_TaxID=10116;
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Pred. No. 2.7e-44;
L; Mismatches 16;
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FAC77FFA82302304
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUCLEOTIDE SEQUENCE.
TISSUE=Spleen;
NIH MGC Project;
Submitted (APR-2005) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EMBL; BC092580; AAH92580.1; -; mRNA.
InterPro; IPR003599; Ig.
InterPro; IPR003597; Ig.
InterPro; IPR003597; Ig.Cl.
InterPro; IPR003596; Ig.WHC.
InterPro; IPR013106; V-set.
Pfam; PF07654; Cl-set; 4.
SMART; SM00409; IG; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG LIKE; 5.
PROSITE; PS00290; IG MHC; UNKNOWN 3.
SEQUENCE 590 AA; 65088 MW; FAC77FF
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                                                                                                             SEQUENCE
Name=RGD1359202;
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                                                                      Sciurognathi
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HV60 MOUSE STANDARD; PRT; 116 AA.
P18531;
01-NOV-1990, integrated into UniProtKB/Swiss-Prot.
01-NOV-1990, sequence version 1.
07-MAR-2006, entry version 39.
Ig heavy chain V region M315 precursor.
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognat Muroidea; Murinae; Mus.
NCBI_TaxID=10090;
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                                                                                                      NUCLEOTIDE SEQUENCE.
STRAIN=BALB/cJ;
MEDLINE=89279149; PubMed=2499654; DOI=10.1084/jem.169.6.2007;
MEDLINE=89279149; PubMed=2499654; DOI=10.1084/jem.169.6.2007;
Levy N.S., Malipiero U.V., Lebecque S.G., Gearhart P.J.;
"Early onset of somatic mutation in immunoglobulin VH genes c primary immune response.";
J. Exp. Med. 169:2007-2019(1989).
                                                                                                                                                                                                                                                                                                  Ig heavy chain V region M315.
/FTId=PRO_0000015241.
Framework-1.
Complementarity-determining-1.
Framework-2.
Complementarity-determining-2.
Framework-3.
By similarity.
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                                                                                                                                                                                                                                                                                    domain; Immunoglobulin V
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Pred. No. 3.4e-44
; Mismatches
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PDB; 1EZV; X-ray; X=20-116.
Ensembl; ENSMUSG0000057048; Mus musculus.
InterPro; IPR007110; Ig-like.
InterPro; IPR013596; Ig_v.
InterPro; IPR013106; V-set.
Pfam; PF07686; V-set; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
3D-structure; Immunoglobulin domain; Immunog SIGNAL
1 18
CHAIN 19 116 Ig heavy chain.
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103
13095 MW;
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Best Local Similarity 87.1%;
Matches 101; Conservative
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Distributed under the Creative Commons Attribution-NoDerive License
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                                                                                 Euteleostomi;
ia; Sciurognathi;
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ia; Sciurognathi;
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MEDLINE=86136012; PubMed=3937730;
Ollier P., Rocca-Serra J., Somme G., Theze J., Fougereau M.;
"The idiotypic network and the internal image: possible regulation a germ-line network by paucigene encoded Ab2 (anti-idiotypic) antibodies in the GAT system.";
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              053VQ5_MOUSE PRELIMINARY; PKI; L.C....
053VQ5;
24-MAY-2005, integrated into UniProtKB/TrEMBL.
24-MAY-2005, sequence version 1.
24-MAY-2006, entry version 4.
07-FEB-2006, entry version 4.
WH-D-JH region (Fragment).
Mus musculus (Mouse).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutelec Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sci Muroidea; Muridae; Musinae; Mus.
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Glires, Rodentia,
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red. No. 8.5e-41;
Mismatches 13
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MEDLINE=86136012; PubMed=3937730;
Ollier P., Rocca-Serra J., Somme G., Theze J.,
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(us musculus (Mouse)

Eukaryota, Metazoa; Chordata; Craniata;

Maimalia; Eutheria; Euarchontoglires; G

Muroldea; Muridae; Murinae; Mus.

NCBI TaxID=10090;
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srPro, IPR003599, Ig.

srPro, IPR007110, Ig-like.

srPro, IPR003596, Ig.v.
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ilarity 81.5%;
Conservative
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IG_LIKE;
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Submitted (NOV-1986)
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SM00406; IGv;
TE; PS50835; IG
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Immunoglobulin C
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Q53VQ1 MOUSE
Q53VQ1;
24-MAY-2005,
27-FEB-2006,
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InterPro;
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MOUSE
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SMART;
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Q53VR3_i
Q53VR3_i
24-MAY-2005, integrated into UniProtKB/TrEMBL.
24-MAY-2005, sequence version 1.
07-FEB-2006, entry version 4.
VH-D-JH region (Fragment).
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi
Muroidea; Muridae; Mus.
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and the internal image: possible regulation
paucigene encoded Ab2 (anti-idiotypic)
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MEDLINE=86136012; PubMed=3937730;
Ollier P., Rocca-Serra J., Somme G., Theze J., Fougereau M.;
"The idiotypic network and the internal image: possible regulation a germ-line network by paucigene encoded Ab2 (anti-idiotypic) antibodies in the GAT system.";
EMBO J. 4:3681-3688(1985).
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Submitted (NOV-1986) to the EMBL/GenBank/DDBJ databases.
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Submitted (NOV-1986) to the EMBL/GenBank/DDBJ databases
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Pred. No. 7.6e-40
3; Mismatches 10
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InterPro; IPR003599; Ig.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
SMART; SM00409; IG; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
Immunoglobulin domain.
NON_TER 115 115
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                       "The idiotypic network and the a germ-line network by paucigen antibodies in the GAT system."; EMBO J. 4:3681-3688(1985).
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13257 MW;
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SMR; Q53VR3; 1-119.
InterPro; IPR003599; Ig.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
SMART; SM00409; IG; 1.
SMART; SM00406; IGv; 1.
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Distributed under the Creative Commons Attribution-NoDerivs License
EMBL; X03375; CAA27077.1; -; mRNA.
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Glires; Rodentia; Sciurognathi;
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MEDLINE=86136012; PubMed=3937730;
Ollier P., Rocca-Serra J., Somme G., Theze J., Fougereau M.;
I'the idiotypic network and the internal image: possible regulation a germ-line network by paucigene encoded Ab2 (anti-idiotypic) antibodies in the GAT system.";
EMBO J. 4:3681-3688(1985).
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Submitted (NOV-1986) to the EMBL/GenBank/DDBJ databases
                                                     Length
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                                       CRC64
                                                                   16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 484.5; DB 2;
Pred. No. 6.7e-39;
; Mismatches 16;
                                       36504D1665BFBB59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      013452306EBAA3BE
                                                    ore 487; DB 2;
ed. No. 3.8e-39;
Mismatches 16
                                                                                                                                                                                   Uni ProtKB/TrEMBL
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Mammalia; Eutheria; Euarchontoglires; (
Muroidea; Muridae; Murinae; Mus.
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EMBL; X03374; CAA27071.1; -; m
SMR; Q53VR7; 1-120.
InterPro; IPR003599; IG.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig-v.
SMART; SM00409; IG; 1.
                             119
13799 MW;
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13892 MW;
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ilarity 79.0%;
Conservative
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07-FEB-2006, ent
VH-D-JH region (
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Q53VQ4;
Q53VQ4;
24-MAY-2005, integrated into UniProtKB/TrEMBL.
24-MAY-2005, sequence version 1.
24-MAY-2006, entry version 3.
VH region (Fragment).
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Murinae; Mus.
NCBI_TaxID=10090;
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi
Muroidea; Muridae; Murinae; Mus.
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MEDLINE=89279149; PubMed=2499654; DOI=10.1084/jem.169.6.2007
Levy N.S., Malipiero U.V., Lebecque S.G., Gearhart P.J.;
"Early onset of somatic mutation in immunoglobulin VH genes
                                                                                                                                                                                                                                                                                                         primary immune response.";
J. Exp. Med. 169:2007-2019(1989).
-!- SIMILARITY: Contains 1 Ig-like (immunoglobulin-like)
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Ig-like.
By similarity.
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Pred. No. 3.9e-38;
5; Mismatches 13;
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                                    UniProtKB/Swiss
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HSSP; P01820; 1G7J.
SMR; P18533; 19-117.
Ensembl; ENSMUSG0000059520; Mus musculus.
InterPro; IPR007110; Ig-like.
InterPro; IPR013106; V-set.
Pfam; PF07686; V-set; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
Immunoglobulin domain; Immunoglobulin V regi
                           01-NOV-1990, integrated into UniProtKJ
01-NOV-1990, sequence version 1.
07-MAR-2006, entry version 37.
Ig heavy chain V region 733 precursor
Mus musculus (Mouse).
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Distributed under the Creative Commons Attribution-NoDerivs License
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Distributed under the Creative Commons Attribution-NoDerivs License
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NUCLEOTIDE SEQUENCE.
MEDLINE=86136012; PubMed=3937730;
Ollier P., Rocca-Serra J., Somme G., Theze J., Fougereau M.;
"The idiotypic network and the internal image: possible regulation a germ-line network by paucigene encoded Ab2 (anti-idiotypic) antibodies in the GAT system.";
EMBO J. 4:3681-3688(1985).
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MEDLINE=86136012; PubMed=3937730;
Ollier P., Rocca-Serra J., Somme G., Theze J., Fougereau M.;
"The idiotypic network and the internal image: possible regulation a germ-line network by paucigene encoded Ab2 (anti-idiotypic) antibodies in the GAT system.";
EMBO J. 4:3681-3688(1985).
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Submitted (NOV-1986) to the EMBL/GenBank/DDBJ databases.
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Submitted (NOV-1986) to the EMBL/GenBank/DDBJ databases
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Pred. No. 6.9e-38
1; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; X03378; CAA27096.1; -; mRNA.

SMR; Q53VQ4; 1-98.

Ensembl; ENSMUSG0000057048; Mus musculus.

InterPro; IPR007110; Ig-like.

InterPro; IPR003596; Ig-v.

SMART; SM00406; IGv; 1.

PROSITE; PS50835; IG_LIKE; 1.

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SMR; Q53VQ9; 1-118.
InterPro; IPR003599; IG.
InterPro; IPR007110; Ig-like
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Best Local Similarity 74.6%; Pred. No. 8e-37;
Matches 88; Conservative 10; Mismatches
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13844 MW;
InterPro; IPR003596; Ig_v.
SMART; SM00409; IG; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1
Immunoglobulin domain.
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US-08-111-080

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US-08-466-163

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US-09-802-077

US-09-940-727

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Sequence 31, Application US/08137117D
Patent No. 5795965
GENERAL INFORMATION:
APPLICANT: TSATO, KON
APPLICANT: SATO, KON
APPLICANT: SATO, KON
APPLICANT: SALDANHA, JOSE
TITLE OF INVENTION: RESERADED HUMAN ANTIBODY TO
TITLE OF INVENTION: RESERVENCES: 158
CCORESPONDENCE ADDRESS:
ADDRESSEE: FOLGY & Lardner
STREED: 20007-5109
CCONTYR: USA
CONTYTER: D.C.
CCONTYR: USA
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OR APPLICATION DATA:
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FILING DATE: 10-OCT-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/462,798
FILING DATE: 05-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Carter, Charles G.
REGISTRATION NUMBER: 35,093
REFERENCE/DOCKET NUMBER: 8648.49
TELECOMMUNICATION INFORMATION:
TELEPHONE: 612/371-5278
TELEPHONE: 612/332-9081
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Patent No. 6111079
GENERAL INFORMATION:
APPLICANT: WYLIE, DWANE E.
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/7
FILING DATE:
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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APPLICANT: MURRAY, PETER
APPLICANT: GOEBEL, PETER
TITLE OF INVENTION: LEAR
TITLE OF SEQUENCES: 46
CORRESPONDENCE ADDRESS:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compati
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CLASSIFICATION: 424
PRIOR APPLICATION DATA:
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; TOPOLOGY: li;
; MOLECULE TYPE:
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STATE: MN
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No. 1.1e-49
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
TELEFAX: (202)672-5399
TELEX: 904136
INFORMATION FOR SEQ ID NO: 31:
SEQUENCE CHARACTERISTICS:
LENGTH: 137 amino
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FILING DATE: 20-DEC-1993
APPLICATION NUMBER: WO PCT/JP92/00544
FILING DATE: 24-APR-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: USA
ZIP: 20007-5109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, V
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/436,717
FILING DATE:
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
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                                                               Score
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FILING DATE: 19-FEB-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 3-95476
FILING DATE: 25-APR-1991
ATTORNEY/AGENT INFORMATION:
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US-08-436-717-31
; Sequence 31, Application US/08436717
; Patent No. 5817790
; GENERAL INFORMATION:
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TYPE: amino TOPOLOGY: 1
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                                                                                                                                                                                                                                                                                                                                                                          Sequence 13, Application US/08308494A
; Patent No. 5959083
; GENERAL INFORMATION:
; APPLICANT: Bosslet, Klaus
APPLICANT: Seeman, Gerhard
; TITLE OF INVENTION: Preparation and Use Thereof
; TITLE OF SEQUENCES: 24
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Finnegan, Henderson, Farabow, Garrett &
; ADDRESSEE: Dunner
                                                                                                                                                                                          Ø
                                                                                                                                                                                          Indel
                                                                                                                                                             Length
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ZIP: 20005-3315
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/308,494A.
FILING DATE: 21-SEP-1994
CLASSIFICATION DATA:
APPLICATION NUMBER: US 07/891,739
FILING DATE: 01-JUN-1992
CLASSIFICATION: 424
                                                                                                                                                           Score 541; DB 2;
Pred. No. 3.4e-45;
; Mismatches 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Kulik, David J.
REGISTRATION NUMBER: 36,576
REFERENCE/DOCKET NUMBER: 05552-1186-02000
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-408-4000
TELEFAX: 202-408-4400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               700
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03-JUN-1991
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                                                                                                                                                         Query Match
Best Local Similarity 87.3%;
Matches 103; Conservative
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INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 117 amino acids
TYPE: amino acid
LENGTH: 119 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: 'linear
                                                                                                   internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
FILING DATE: 03-JUN-
                                                       MOLECULE TYPE: protein HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS: si
TOPOLOGY: linear
                                                                                              FRAGMENT TYPE:

ORIGINAL SOURCE:
US-08-767-128-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: 1 MOLECULE TYPE:
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US-08-308-494A-13
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                                           TOPOLOGY:
                                                                                     ANTI-SENSE:
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  Length
                       Indels
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APPLICANT: SEINO, Ken-ichiro
APPLICANT: KAYAGAKI, No. 6068841uhiko
APPLICANT: YAGITA, Hideo
APPLICANT: VAGITA, Ko
APPLICANT: OKUMURA, Ko
APPLICANT: NAKATA, Motomi
TITLE OF INVENTION: THERAPEUTIC AGENT FOR HEPATITIS
NUMBER OF SEQUENCES: 18
CORRESPONDENCE ADDRESS:
ADDRESSEE: McDermott, Will & Emery
STREET: 99 Canal Center Plaza
CITY: Alexandria
STATE: Virginia
COUNTRY: USA
ZIP: 22314
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 2; L.1.4e-42;
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             .1e-43
  DB
Score 525.5;
Pred. No. 1.1e
7; Mismatches
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Pred. No. 1.4e
8; Mismatches
                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/065,059
FILING DATE:
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Bucca Ph.D., Daniel
REGISTRATION NUMBER: P-42,368
REFERENCE/DOCKET NUMBER: 50356-151
TELEPHONE: 703-518-5100
TELEFAX: 703-684-1124
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                    A:
US/09/065,059
                                                                                                                                                                  11, Application US/09065059
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71.3%;
ilarity 85.2%;
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Similarity 83.1%;
98; Conservative
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         l Similarity
98; Conserv
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US-09-065-059-11
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Query Match
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Matches 98
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Best Local S
Matches 98
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Patent No.
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OPERATING
SOFTWARE:
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Best Local S
Matches 98
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Sequence 11, Application US/08913555
Patent No. 6946255
GENERAL INFORMATION:
APPLICANT: KAYAGAKI, No. 6946255uhiko
APPLICANT: YAGITA, Kideo
APPLICANT: OKUMURA, Ko
APPLICANT: OKUMURA, MOLOMI
TITLE OF INVENTION: MONOCLONAL ANTIBODY SPECIFICALLY
TITLE OF INVENTION: REACTING WITH FAB LIGAND AND PRODUCTION PROCESS
NUMBER OF SEQUENCES: 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                   COUNTRY: COUNTRY: COUNTRY: COUNTRY: COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
COMPUTER: BATENTIN PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/913,555
FILING DATE: 19-SEP-1997
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: BUCCA Ph.D., Daniel
REGISTRATION NUMBER: 42,368
REFERENCE/DOCKET NUMBER: 50356-150
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-756-8609
"PELECOMMUNICATION INFORMATION:
TELEPHONE: 202-756-8609
"PELECOMMUNICATION OF 11:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 514; DB 2;
Pred. No. 1.4e-42
}; Mismatches
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; Sequence 23, Application 08/111080
; Patent No. 5558865
; GENERAL INFORMATION:
; APPLICANT: Ohno, Tsuneya
; TITLE OF INVENTION: HIV Immunotherapeutics
; NUMBER OF SEQUENCES: 38
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Marshall, O'Toole, Gerstein, Muraddonessee: Borun
; STREET: 6300 Sears Tower, 233 S. Wacker Dri
                                                                                                                                                                               Emery
1, Suite 300
                                                                                                                                                       STREET: 99 Canal Center Plaza, CITY: Alexandria STATE: Viroinia
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Similarity 83.1%;
98; Conservative
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 118 amino acids
TYPE: amino acid
                                                                                                                                           NUMBER OF SEQUENCES: 31
CORRESPONDENCE ADDRESS:
ADDRESSEE: McDermott,
STREET: 99 Canal Cent
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: USA
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STATE: Illinois
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US-08-913-
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                                  Version #1.2
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; Sequence 23, Application US/08211980
; Patent No. 5665569
; GENERAL INFORMATION:
; TITLE OF INVENTION: HIV Immunotherapeutics
; VUMBER OF SEQUENCES: 38
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Marshall, O'Toole, Gerstein, Muraphores and Street: 6300 Sears Tower, 233 S. Wacker Drices and Street: Illinois
; CITY: Chicago
; CITY: USA
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OB
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                                                                                                 CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/748,562
FILING DATE: 22-AUG-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/07111
FILING DATE: 24-AUG-1992
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                      APPLICATION NUMBER: US 08/039,457
FILING DATE: 22-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Borun, Michael F.
REGISTRATION NUMBER: 25,447
REFERENCE/DOCKET NUMBER: 31629
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MBER: PCT/US92/07111
24-AUG-1992
 compatible
PC-DOS/MS-DOS
In Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Patentin Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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US/08/211,
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474-0448
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Similarity 83.1%;
98; Conservative
                                                                                                                                                                                                                                                                                                                                                                 TELEX: 25-3856
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 114 amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
FILING DATE:
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT
COMPUTER: IBM PC comp
OPERATING SYSTEM: PC-
SOFTWARE: Patentin Re
CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  protein
                                                           APPLICATION NUMBER:
FILING DATE:
CLASSIFICATION: 42
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US-08-111-080-23
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TYPE:
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Matches
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/07967
FILING DATE:
CLASSIFICATION DATA:
APPLICATION NUMBER: PCT/US92/07111
FILING DATE: 24-AUG-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/039,457
FILING DATE: 22-APR-1993
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 9
PCT-US93-07967-23
; Sequence 23, Application PC/TUS9307967
; GENERAL INFORMATION:
; APPLICANT: Ohno, Tsuneya
; TITLE OF INVENTION: HIV Immunotherap;
; NUMBER OF SEQUENCES: 38
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Marshall, O'Toole, Gers
; ADDRESSEE: Borun
; STREET: 6300 Sears Tower, 233 S. W
; CITY: Chicago
; STATE: Illinois
; COUNTRY: USA
         APPLICATION NUMBER: US 08/039,457
FILING DATE: 22-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Borun, Michael F.
REGISTRATION NUMBER: 25,447
REFERENCE/DOCKET NUMBER: 31629
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 474-6300
TELEFAX: (312) 474-0448
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 31629
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REGISTRATION NUMBER: 25,447
REFERENCE/DOCKET NUMBER: 3162
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 474-6300
TELEFAX: (312) 474-648
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Similarity 83.1%;
98; Conservative
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nino acid
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                                                                                                                           IELEX: 25-3856
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 114 aming
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TELEX: 25-3856
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 114 aming
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APPLICATION DATA:
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No. 4.6e-42;
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                                                      ore 510.5; DB ed. No. 3e-42; Mismatches
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MEDIUM TYPE: 3.5 inch, 1.44 Mb fl.

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: WinPatin (Genentech)

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/466,151

FILING DATE:
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US/08/466,151
                                                       Score
Pred.
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Pred.
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/87940F
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07-MAY-1992
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14-AUG-1991
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APPLICATION NUMBER: 08/466163
FILING DATE: 06-Jun-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 08/405617
FILING DATE: 15-MAR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/185899
                                                                                                                                                                                                                                                           7, Application US/08466151
5. 6037453
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Svoboda, Craig G.
REGISTRATION NUMBER: 39,044
REFERENCE/DOCKET NUMBER: P071
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1489
TELEFAX: 650/952-9881
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Craig G.
----- 39,044
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Patent No. 6037453
GENERAL INFORMATION:
APPLICANT: Jardieu, Paula M.
TITLE OF INVENTION: Immunoglol
NUMBER OF SEQUENCES: 65
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
STREET: 1 DNA Way
CITY: South San Francisco
STATE: California
COUNTRY: USA
ZIP: 94080
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ilarity 83.1%;
Conservative
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81.0%;
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/7
FILING DATE: 14-AUG-1991
ATTORNEY/AGENT INFORMATION:
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CHARACTERISTICS
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TOPOLOGY: 1 MOLECULE TYPE: PCT-US93-07967-23
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TELEFAX: 65
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US-08-466-151-7
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Sequence 7, Application US/09802077

Batent No. 6699472

GENERAL INFORMATION:

APPLICANT: Jardieu, Paula M.

APPLICANT: Jardieu, Paula M.

TITLE OF INVENTION: Method of Treating Allergic Disorders (as FILE REFERENCE: P0718P2C2US

CURRENT APPLICATION NUMBER: US/09/802,077

CURRENT FILING DATE: 2001-03-08

PRIOR APPLICATION NUMBER: US 08/185,899

PRIOR APPLICATION NUMBER: US 08/185,899

PRIOR FILING DATE: 1994-01-26

PRIOR FILING DATE: 1992-08-14

PRIOR APPLICATION NUMBER: US 07/879,495

PRIOR APPLICATION NUMBER: US 07/879,495
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Pred. No. 4.6e-42;
; Mismatches 13
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Pred. No. 4.6e-42
7; Mismatches 13
PRIOR APPLICATION NUMBER: PCT/US92/06860
PRIOR FILING DATE: 1992-08-14
PRIOR APPLICATION NUMBER: US 07/879,495
PRIOR FILING DATE: 1992-05-07
PRIOR APPLICATION NUMBER: US 07/744,768
PRIOR FILING DATE: 1991-08-14
NUMBER OF SEQ ID NOS: 64
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ilarity 81.0%;
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PRIOR APPLICATION NUMBER: US (
PRIOR FILING DATE: 1991-08-14
NUMBER OF SEQ ID NOS: 64
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Similarity 81.0%;
98; Conservative
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LENGTH: 10
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                                          DVQHQESEPDLVKPSQSLSLTCTVTGYSITSGYNRHWIRQFPGNKLEWMGYIHYSGSTNY
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Pred. No. 4.6e-42;
; Mismatches 13;
                                                                                                                                                                                                                     Sequence 7, Application US/08466163B
Fatent No. 6329509
GENERAL INFORMATION:
APPLICANT: Jardieu, Paula M.
APPLICANT: Presta, Leonard G.
TITLE OF INVENTION: Immunoglobulin Variants
FILE REFERENCE: P0718P2C1D1
CURRENT APPLICATION NUMBER: US/08/466,163B
CURRENT FILING DATE: 1995-06-06
PRIOR APPLICATION NUMBER: US 08/185,899
PRIOR FILING DATE: 1994-01-26
PRIOR FILING DATE: 1994-01-26
PRIOR FILING DATE: 1992-05-07
PRIOR FILING DATE: 1992-05-07
PRIOR FILING DATE: 1991-08-14
NUMBER OF SEQ ID NOS: 64
    Mismatches
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Patent No. 6685939
GENERAL INFORMATION:
"APPLICANT: Jardieu, Paula M.
"NPPLICANT: Presta, Leonard G.
"ITLE OF INVENTION: Method of Preventing :ILE REFERENCE: P0718P2C3US
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CURRENT FILING DATE: 2001-03-08

PRIOR APPLICATION NUMBER: US 08/405,617

PRIOR FILING DATE: 1995-03-15

PRIOR APPLICATION NUMBER: US 08/185,899

PRIOR FILING DATE: 1994-01-26
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-09-802-096-7
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Pred. No. 4.6e-42;
7; Mismatches 13;
RESULT 14
US-09-925-179-7
US-09-925-179-7

Sequence 7, Application US/09925179

Fatent No. 6914129

GENERAL INFORMATION:

APPLICANT: Jardieu, Paula M.

TITLE OF INVENTION: Anti-IgE Antibodies (as ame FILE REFERENCE: P0718P2C1D1C1US

CURRENT APPLICATION NUMBER: US/09/925,179

CURRENT FILING DATE: 2001-08-08

PRIOR APPLICATION NUMBER: US 08/466,163

PRIOR APPLICATION NUMBER: US 08/466,163

PRIOR PILING DATE: 1995-06-06

PRIOR PILING DATE: 1995-03-15

PRIOR PILING DATE: 1995-03-15

PRIOR PILING DATE: 1995-01-26

PRIOR FILING DATE: 1992-08-14

PRIOR PILING DATE: 1992-08-14

PRIOR FILING DATE: 1992-08-14

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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/672,345C
FILING DATE: 24-JUN-1996
CLASSIFICATION: 435
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TITLE OF INVENTION: ANTI-COCAINE CANUMBER OF SEQUENCES: 108
CORRESPONDENCE ADDRESS:
ADDRESSEE: Cooper and Dunham LLP
STREET: 1185 Avenue of the Americ
CITY: New York
STATE: New York
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 13, Application US/08672345C
Patent No. 5948658
GENERAL INFORMATION:
APPLICANT: Landry Donald, W.
TITLE OF INVENTION: ANTI-COCAINE
NUMBER OF SEQUENCES: 108
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PRIOR APPLICATION NUMBER: US (
PRIOR FILING DATE: 1991-08-14
NUMBER OF SEQ ID NOS: 68
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Similarity 81.0%;
98; Conservative
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ORGANISM: Mus musculus
19-925-179-7
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Pred. No. 4.3e-42
; Mismatches 10
       NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 0575/51400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 212-278-0400
TELEFAX: 212-391-0525
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 117 amino acids
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nilarity 82.2%;
Conservative
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117 amino acids

STRANDEDNESS: single

; TOPOLOGY: linear

; MOLECULE TYPE: proteir

US-08-672-345C-13
INFORMATION:
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ne : 40.3968 secs
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Pred. No. 2e-49;
6; Mismatches
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Publication No. US20050152907A1
GENERAL INFORMATION:
APPLICANT: LIANG, Tony W.
APPLICANT: LOO, Deryk T.
APPLICANT: XU, Xiaolin
TITLE OF INVENTION: KID3 AND KID3 ANTIBODIES
TITLE OF INVENTION: THERETO
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US-09-791-551-109
; Sequence 109, Application US/09791551
; Publication No. US20030235584A1
; GENERAL INFORMATION:
; APPLICANT: KLOETZER, WILLIAM S.
; APPLICANT: HANNA, NABIL
; TITLE OF INVENTION: METHOD FOR PREPARING ANT
; FILE REFERENCE: 037003/0277869
; CURRENT APPLICATION NUMBER: US/09/791,551
; CURRENT FILING DATE: 2001-02-26
; PRIOR FILING DATE: 2000-02-28
; PRIOR FILING DATE: 2000-02-28
; PRIOR FILING DATE: 2000-09-18
; NUMBER OF SEQ ID NOS: 119
; SOFTWARE: Patentin Ver: 2.1
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/837,904
FILING DATE: 04-May-2004
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                         10;
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FILING DATE: 13-Jul-1998
APPLICATION NUMBER: US 08/436,717
FILING DATE: 08-MAY-1995
APPLICATION NUMBER: US 08/137,117
FILING DATE: 20-DEC-1993
APPLICATION NUMBER: WO PCT/JP92/00544
FILING DATE: 24-APR-1992
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RECEPTOR
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BENDIG, Mary Margaret
JONES, Steven Tarran
SALDANHA, Jose William
F INVENTION: RESHAPED HUMAN AL
FILE REFERENCE: 415072002700
CURRENT APPLICATION NUMBER: US/10/943,640
CURRENT FILING DATE: 2004-09-17
PRIOR APPLICATION NUMBER: US 60/504,441
PRIOR FILING DATE: 2003-09-18
NUMBER OF SEQ ID NOS: 4
SOFTWARE: FastSEQ for Windows Version 4.0
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COUNTRY: U.S.A.
ZIP: 20007-5109
COMPUTER READABLE FORM:
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US-10-837-904-31
; Sequence 31, Application US/10837904
; Publication No. US20050142635A1
; GENERAL INFORMATION:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: FOLEY & LA
STREET: 3000 K Street
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FILING DATE: 19-FEB
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illarity 87.5%;
Conservative
                                                                                                                                                          Synthetic
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nes 119; Conser
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OTHER INFORMATION:
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CITY: Wa
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                                                                           SOFTWARE: Fas
SEQ ID NO 4
LENGTH: 468
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; APPLICANT: HANNA, NABIL; APPLICANT: HANNA, NABIL; TITLE OF INVENTION: METHOD FOR PREPARING ANTI-MIF ANTIBODIES; FILE REFERENCE: 037003/0277869; CURRENT APPLICATION NUMBER: US/09/791,551; CURRENT FILING DATE: 2001-02-26; PRIOR APPLICATION NUMBER: 60/185,390; PRIOR FILING DATE: 2000-02-28; PRIOR FILING DATE: 2000-09-18; PRIOR FILING DATE: 2000-09-18
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1.8e-43;
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Pred. No. 1.2e
8; Mismatches
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No. 1
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FILING DATE: 25-APR-1991

ATTORNEY/AGENT INFORMATION:
NAME: Wegner, Harold C.
REGISTRATION NUMBER: 25,258
REFERENCE/DOCKET NUMBER: 53,
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 672-5399
TELEFAX: (202) 672-5399
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ilarity 78.4%;
Conservative
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PatentIn Ver. 2
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: Local Similarity
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114; Conser
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Q ID NO 36
LENGTH: 121
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Best Local S
Matches 105
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Best Local S
Matches 103
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US-11-012-353
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APPLICANT: CORVAIA, NATHALIE
APPLICANT: DUFLOS, ALAIN
APPLICANT: HAEUW, JEAN-FRANCOIS
APPLICANT: LEGER, OLIVIER
APPLICANT: BECK, ALAIN
TITLE OF INVENTION: NOVEL ANTI-IGF-IR AND/OR ANTI-INSULIN/IGF-I
TITLE OF INVENTION: RECEPTORS ANTIBODIES AND USES THEREOF
FILE REFERENCE: 017753-198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 127
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Pred. No. 1.4e-42
; Mismatches 1.
                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/735,916A CURRENT FILING DATE: 2003-12-16
PRIOR APPLICATION NUMBER: FR 03/08 538
PRIOR FILING DATE: 2003-07-11
PRIOR APPLICATION NUMBER: PCT/FR 03/00 178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/11/012,353
CURRENT FILING DATE: 2004-12-16
PRIOR APPLICATION NUMBER: 10/735,916
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APPLICATION NUMBER: FR 0308538
FILING DATE: 2003-07-11
APPLICATION NUMBER: PCT/FR03/00178
RESULT 5
US-10-735-916A-52
Sequence 52, Application US/10735916A
Publication No. US20050084906A1
GENERAL INFORMATION:
APPLICANT: GOETSCH, Liliane
APPLICANT: CORVAIA, Nathalie
APPLICANT: LEGER, Olivier
APPLICANT: BECK, Alain
APPLICANT: BECK, Alain
                                                                                           APPLICANT: CORVAIA, Nathalie
APPLICANT: LEGER, Olivier
APPLICANT: DUFLOS, Alain
APPLICANT: BECK, Alain
APPLICANT: HAEUW, Jean-Francois
TITLE OF INVENTION: NOVEL ANTI-IGF-IR
FILE REFERENCE: 017753-183
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US-11-012-353-52

Sequence 52, Application US/11012353

Publication No. US20050249730A1

GENERAL INFORMATION:
APPLICANT: GOETSCH, LILIANE
APPLICANT: CORVAIA, NATHALIE
APPLICANT: DUFLOS, ALAIN
APPLICANT: HAEUW, JEAN-FRANCOIS
APPLICANT: LEGER, OLIVIER
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R APPLICATION NUMBER: FR 02
R FILING DATE: 2002-01-18
R APPLICATION NUMBER: FR 02
R FILING DATE: 2002-01-18
R APPLICATION NUMBER: FR 02
R FILING DATE: 2002-05-07
ER OF SEQ ID NOS: 156
WARE: Patentin Ver. 2.1
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ilarity 82.0%;
Conservative
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nes 105; Conser
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ENGTH: 127
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Pred. No. 1.1e-41;
1; Mismatches 11;
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Pred. No. 1.4e-42;
5; Mismatches 13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Containing
                                                                                                                                                                                                                                                                                                                                                                                                                        US-10-310-674A-36
; Sequence 36, Application US/10310674A
; Publication No. US20030166860A1
; GENERAL INFORMATION:
; APPLICANT: TeGenero GmbH
; TITLE OF INVENTION: Family
; TITLE OF INVENTION: Family
; FILE REFERENCE: 00140/004001
; CURRENT APPLICATION NUMBER: US/10/310,674A
; CURRENT FILING DATE: 2002-12-04
; NUMBER OF SEQ ID NOS: 44
; SOFTWARE: Patentin version 3.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: No. US20030166860Alel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-10-389-679-12; Sequence 12, Application US/10389679; Publication No. US20040092718A1; GENERAL INFORMATION:
; APPLICANT: TeGenero AG; APPLICANT: TeGenero AG; APPLICANT: HUNIG, Thomas
    FR 0205753
                             0200653
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PRIOR APPLICATION NUMBER: FR 020 PRIOR FILING DATE: 2002-05-07 PRIOR APPLICATION NUMBER: FR 020 PRIOR FILING DATE: 2002-01-18 PRIOR FILING DATE: 2002-01-18 NUMBER OF SEQ ID NOS: 162 SOFTWARE: Patentin Ver. 3.3 SEQ ID NO 52 LENGTH: 127 TYPE: PRT
                                                                                                                                                                                   ch 75.2%;
1 Similarity 82.0%;
105; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ch 73.8%;
il Similarity 87.3%;
103; Conservative
                                                                                                                                             Mus musculus
-52
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WGQGTTLT
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APPLICANT: Fandl, James
APPLICANT: Chen, Gang
APPLICANT: Papadopoulos, Nicholas
APPLICANT: Aldrich, Thomas F.
TITLE OF INVENTION: Fusion Polypeptides CarlLE REFERENCE: 1080A
CURRENT APPLICATION NUMBER: US/11/035,599
CURRENT FILING DATE: 2005-01-14
PRIOR APPLICATION NUMBER: 60/536,968
PRIOR FILING DATE: 2004-01-16
NUMBER OF SEQ ID NOS: 53
SOFTWARE: FastSEQ for Windows Version 4.0
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Pred. No.
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ilarity 87.3%;
Conservative
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ilarity 84.0%;
Conservative 1
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les 100; Conser
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LENGTH: 730
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LENGTH: 121
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No. 1.1e-41
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Pred. No. 1.1e-41
4; Mismatches 11
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Pred. No. 1.1e
4; Mismatches
TITLE OF INVENTION: Use of a CD28 Binding Strint OF INVENTION: Composition
FILE REFERENCE: 00140/009001, TEG/US/0302
CURRENT APPLICATION NUMBER: US/10/389,679
CURRENT FILING DATE: 2003-03-13
PRIOR FILING DATE: 2002-03-13
NUMBER OF SEQ ID NOS: 19
SOFTWARE: Patentin version 3.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 12, Application US/10988207

Publication No. US20060008457A1

GENERAL INFORMATION:

APPLICANT: TeGenero AG

TITLE OF INVENTION: USE OF AN EFFECTIVE SUBS

TITLE OF INVENTION: PHARMACEUTICAL COMPOSIT

FILE REFERENCE: TEG/US/0409

CURRENT APPLICATION NUMBER: US/10/988,207

CURRENT FILING DATE: 2004-11-12

PRIOR APPLICATION NUMBER: DE 103 52 900.4

PRIOR FILING DATE: 2003-11-11
    a CD28 Binding
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Sequence 12, Application US/10946836A
Publication No. US20060009382A1
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ilarity 87.3%;
Conservative
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11 Similarity 87.3%;
103; Conservative
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SOFTWARE: Patentin version
SEQ ID NO 12
LENGTH: 121
                                                                                                                                                                             FEATURE:

NAME/KEY: misc feature:

LOCATION: (1)..(121)

OTHER INFORMATION: mab
US-10-389-679-12
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US-10-988-207-12
                                                                                                                                                  TYPE: PRT
ORGANISM: artificial
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ORGANISM: Artificial
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LENGTH: 121
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US-10-946-
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APPLICANT: Tegenero AG

APPLICANT: Hanke, Thomas
APPLICANT: Hanke, Thomas
APPLICANT: Lin, Chia-Huey
TITLE OF INVENTION: USE OF A CD28 BINDING PHARMACEUTICAL SUBSTANCE FOR MAK:
TITLE OF INVENTION: PHARMACEUTICAL COMPOSITION WITH DOSIS DEPENDENT EFFECT
FILE REFERENCE: 7003/25, TEG/US/0407
CURRENT APPLICATION NUMBER: US/10/946,836A
CURRENT FILING DATE: 2004-09-22
PRIOR FILING DATE: 2003-09-22
PRIOR FILING DATE: 2003-10-20
PRIOR FILING DATE: 2003-10-20
NUMBER OF SEQ ID NOS: 19
SOFTWARE: Patentin version 3.2
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1.1e-41;
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Pred. No. 1.9e-40;
11; Mismatches 5
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                                                      Sequence 31, Application US/11035599; Publication No. US20050158829A1; GENERAL INFORMATION:
APPLICANT: Fandl, James
APPLICANT: Chen, Gang
APPLICANT: Papadopoulos, Nicholas
APPLICANT: Aldrich, Thomas F.
TITLE OF INVENTION: Fusion Polypeptides Capa; TITLE OF INVENTION: Activating Receptors
FILE REFERENCE: 1080A
CURRENT APPLICATION NUMBER: 60/536,968; PRIOR APPLICATION NUMBER: 60/536,968; PRIOR FILING DATE: 2004-01-16
NUMBER OF SEQ ID NOS: 53
SOFTWARE: FastSEQ for Windows Version 4.0
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APPLICANT: Chen, Gang
APPLICANT: Chen, Gang
APPLICANT: Papadopoulos, Nicholas
APPLICANT: Aldrich, Thomas F.
TITLE OF INVENTION: Fusion Polypeptides
TITLE OF INVENTION: Activating Receptor
FILE REFERENCE: 1080A
CURRENT APPLICATION NUMBER: US/11/035,59;
CURRENT FILING DATE: 2005-01-14
PRIOR APPLICATION NUMBER: 60/536,968
PRIOR FILING DATE: 2004-01-16
NUMBER OF SEQ ID NOS: 53
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 29
LENGTH: 761
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US-11-035-599-29
Sequence 29, Application US/1103559
Publication No. US20050158829A1
GENERAL INFORMATION:
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Best Local Similarity 84.0%;
Matches 100; Conservative 11
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ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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                                             US-11-035-599-31
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; Sequence 53, Application US/09874141
; Publication No. US20030012781A1
; GENERAL INFORMATION:
; APPLICANT: ANDERSON, DARRELL
; APPLICANT: HANNA, NABIL
; APPLICANT: KLOETZER, WILLIAM H.
; APPLICANT: KLOETZER, WILLIAM S.
; TITLE OF INVENTION: NON-AGONISTIC ANTIBODIES TO HUMAN GP39, COMF
; TITLE OF INVENTION: CONTAINING, AND THERAPEUTIC USE THEREOF
; TITLE OF INVENTION NUMBER: US/09/874,141
; CURRENT APPLICATION NUMBER: 60/209,584
; PRIOR FILING DATE: 2000-06-06
; PRIOR FILING DATE: 2000-06-06
; NUMBER OF SEQ ID NOS: 53
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US-11-035-599-28
is Sequence 28, Application US/11035599
j Sequence 28, Application US/11035599
j Publication No. US20050158829A1
j GENERAL INFORMATION:
j APPLICANT: Fandl, James
j APPLICANT: Chen, Gang
j APPLICANT: Papadopoulos, Nicholas
j APPLICANT: Aldrich, Thomas F.
j TITLE OF INVENTION: Fusion Polypeptides Capable of
j TITLE OF INVENTION: Activating Receptors
j FILE REFERENCE: 1080A
j CURRENT APPLICATION NUMBER: 60/536,968
j CURRENT FILING DATE: 2004-01-14
j PRIOR FILING DATE: 2004-01-16
j NUMBER OF SEQ ID NOS: 53
j SOFTWARE: FastSEQ for Windows Version 4.0
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ORGANISM: Artificial Sequence
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Best Local Similarity 84.0%;
Matches 100; Conservative
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76.5%;
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US-09-874-141-53
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; OTHER INFORMATION:
US-11-035-599-28
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US-09-874-141-53
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LENGTH: 762
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Indels

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12; Mismatches

104; Conservative

Matches

Search completed: May 19, 2006, 17:52:56 Job time : 134.841 secs .rapbn

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2/pubpaa/US06 NEW PUB.pep: *
2/pubpaa/US07 NEW PUB.pep: *
2/pubpaa/US08 NEW PUB.pep: *
2/pubpaa/PCT NEW PUB.pep: *
2/pubpaa/US10 NEW PUB.pep: *
2/pubpaa/US11 NEW PUB.pep: *
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Sequence 11, Application US/11219121

Publication No. US20060093601A1

GENERAL INFORMATION:

APPLICANT: Fong, Sherman

APPLICANT: Pong, Sherman

TITLE OF INVENTION: HUMANIZED ANTI-BETA7 ANTAGONISTS AND

FILE REFERENCE: P2159R1

CURRENT APPLICATION NUMBER: US/11/219,121

CURRENT FILING DATE: 2005-09-02

PRIOR FILING DATE: 2004-09-03

NUMBER OF SEQ ID NOS: 68

SEQ ID NO 11

LENGTH: 117

TYPE: PRT

ORGANISM: Artificial sequence

FEATURE:

ORGANISM: Artificial sequence is synthesized
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; Publication No. US20060093601A1
; GENERAL INFORMATION:
; APPLICANT: Fong, Sherman
; APPLICANT: Dennis Mark S.
; TITLE OF INVENTION: HUMANIZED ANTI-BETA7 ANTAGONISTS ANI
; FILE REFERENCE: P2159R1
; CURRENT APPLICATION NUMBER: US/11/219,121
; CURRENT FILING DATE: 2005-09-02
; PRIOR APPLICATION NUMBER: US 60/607,377
; PRIOR FILING DATE: 2004-09-03
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Pred. No. 8.5e-26;
15; Mismatches 27
                                                                                                      Score 491; DB 7;
Pred. No. 8.1e-31
8; Mismatches 13
                                                                                                                                                                                                                                                                             Sequence 30, Application US/11211917

Sequence 30, Application US/11211917

Publication No. US20060093600A1

GENERAL INFORMATION:

APPLICANT: BEDIAN, VAHE

APPLICANT: GLADUE, RONALD P.

APPLICANT: CORVALAN, JOSE

APPLICANT: JIA, XIAO-CHI

APPLICANT: FENG, XIAO

TITLE OF INVENTION: ANTIBODIES TO CD40

FILE REFERENCE: ABX-PF/3 US

CURRENT APPLICATION NUMBER: US/11/211,917

CURRENT FILING DATE: 2005-08-25

PRIOR APPLICATION NUMBER: 60/348,980

PRIOR FILING DATE: 2001-11-08

PRIOR FILING DATE: 2001-11-09

NUMBER OF SEQ ID NOS: 147

SOFTWARE: Patentin Ver. 2.1
                                                                      sequence is synthesized
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-11-211-917-70
; Sequence 70, Application US/11211917
; Publication No. US20060093600A1
; CENERAL INFORMATION:
; APPLICANT: BEDIAN, VAHE
; APPLICANT: GLADUE, RONALD P.
; APPLICANT: CORVALAN, JOSE
                                                                                                     66.6%;
illarity 79.7%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                 TYPE: PRT
ORGANISM: Artificial &
FEATURE:
OTHER INFORMATION: SEC
 NOS:
                                                                                                                  Similarity 94; Conser
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 SEQ
                                                                                                                                                                                                                                                            RESULT 3
US-11-211-917-30
NUMBER OF SEQ
EQ ID NO 13
LENGTH: 146
                                                                                US-11-219-121-13
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Best Local S
Matches 87
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Best Lical
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Pred. No. 8.5e-26;
5; Mismatches 24
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APPLICANT: JIA, XIAO-CHI
APPLICANT: FENG, XIAO
TITLE OF INVENTION: ANTIBODIES TO CD40
FILE REFERENCE: ABX-PF/3 US
CURRENT APPLICATION NUMBER: US/11/211,917
CURRENT FILING DATE: 2005-08-25
PRIOR APPLICATION NUMBER: US/10/292,088
PRIOR FILING DATE: 2002-11-08
PRIOR APPLICATION NUMBER: 60/348,980
PRIOR FILING DATE: 2001-11-09
NUMBER OF SEQ ID NOS: 147
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 70
LENGTH: 466
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08-25
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; Publication No. US20060093600A1
; GENERAL INFORMATION:
; APPLICANT: BEDIAN, VAHE
; APPLICANT: GLADUE, RONALD P.
APPLICANT: GLADUE, RONALD P.
APPLICANT: CORVALAN, JOSE
; APPLICANT: FENG, XIAO
; TITLE OF INVENTION: ANTIBODIES TO CD40
; FILE REFERENCE: ABX-PF/3 US
; CURRENT APPLICATION NUMBER: US/11/211,917
; CURRENT FILING DATE: 2005-08-25
; PRIOR FILING DATE: 2002-11-08
; PRIOR FILING DATE: 2001-11-09
; NUMBER OF SEQ ID NOS: 147
; SOFTWARE: PATENTIN Ver. 2.1
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ilarity 64.7%;
Conservative 1
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65.4%;
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                                                                                                                                                                                                                                            Homo sapiens
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US-11-211-917-86
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US-10-981-300-20
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LENGTH: 118
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Best Local S
Matches 80
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US-10-981-300
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Matches
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US-10-981-300-51
; Sequence 51, Application US/10981300
; Publication No. US20060093599A1
; GENERAL INFORMATION:
; APPLICANT: GADI GAZIT-BORNSTEIN
; TITLE OF INVENTION: FOR MAKING AND USING THE SAM
; FILE REFERENCE: ABGX-005
; CURRENT APPLICATION NUMBER: US/10/981,300
; CURRENT FILING DATE: 2004-11-03
; CURRENT FILING DATE: 2004-11-03
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1.4e-25;
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                                                                                                                                APPLICANT: BEDIAN, VAHE
APPLICANT: GLADUE, RONALD P.
APPLICANT: GLADUE, RONALD P.
APPLICANT: CORVALAN, JOSE
APPLICANT: JIA, XIAO-CHI
APPLICANT: FENG, XIAO
TITLE OF INVENTION: ANTIBODIES TO CD40
FILE REFERENCE: ABX-PF/3 US
CURRENT APPLICATION NUMBER: US/11/211,917
CURRENT FILING DATE: 2005-08-25
PRIOR APPLICATION NUMBER: US/10/292,088
PRIOR APPLICATION NUMBER: 60/348,980
                                                                                                                                                                                                                                                                                                                                                                                                                             Score 419.5;
Pred. No. 1.
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; Pred. No. 1
13; Mismatch
                                                                                          Sequence 142, Application US/11211917
Publication No. US20060093600A1
GENERAL INFORMATION:
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ilarity 69.5%;
Conservative 1
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70.1%;
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NUMBER OF SEQ ID NOS: 147
SOFTWARE: Patentin Ver. 2.1
TTYGYFDVWGQGTTVT
                       -DYGWFAPWGQGTLVT
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82; Conserv
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82; Conser
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Best Local S
Matches 82
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Matches 82
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Sequence 20, Application US/10981300
; Publication No. US20060093599A1
; GENERAL INFORMATION:
; APPLICANT: GIORGIO SENALDI
; APPLICANT: GADI GAZIT-BORNSTEIN
; TITLE OF INVENTION: ANTI-PROPERDIN ANTIBODIES, AN
; TITLE OF INVENTION: FOR MAKING AND USING THE SAM
; FILE REFERENCE: ABGX-005
; CURRENT APPLICATION NUMBER: US/10/981,300
; CURRENT FILING DATE: 2004-11-03
; NUMBER OF SEQ ID NOS: 71
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 20
; LENGTH: 443
                                                                                            Sequence 54, Application US/10981300
; Publication No. US20060093599A1
; GENERAL INFORMATION:
; APPLICANT: GIORGIO SENALDI
; APPLICANT: GADI GAZIT-BORNSTEIN
; TITLE OF INVENTION: ANTI-PROPERDIN ANTIBODIES, AN
; TITLE OF INVENTION: FOR MAKING AND USING THE SAM
; FILE REFERENCE: ABGX-005
; CURRENT APPLICATION NUMBER: US/10/981,300
; CURRENT FILING DATE: 2004-11-03
; NUMBER OF SEQ ID NOS: 71
; SOFTWARE: FastSEQ for Windows Version 4.0
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Pred. No. 5.3e-25
16; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                Score 418; DB 6;
Pred. No. 1.6e-25
5; Mismatches 1
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Sequence 109, Application US/11211
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80; Conservative
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US-10-981-300-54
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ID NO 66
ENGTH: 121
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ORGANISM:
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Best Local S
Matches 82
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Best Local S
Matches 82
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Pred. No. 4.6e-25;
4; Mismatches 22;
                                                                                                                                                                                                                                                                                                                                Score 417.5; DB 7;
Pred. No. 1.8e-25;
15; Mismatches 17;
                                         APPLICANT: GLADUE, RONALD P.
APPLICANT: CORVALAN, JOSE
APPLICANT: JIA, XIAO-CHI
APPLICANT: JIA, XIAO-CHI
APPLICANT: FENG, XIAO
TITLE OF INVENTION: ANTIBODIES TO CD40
FILE REFERENCE: ABX-PF/3 US
CURRENT APPLICATION NUMBER: US/11/211,917
CURRENT FILING DATE: 2005-08-25
PRIOR APPLICATION NUMBER: US/10/292,088
PRIOR FILING DATE: 2002-11-08
PRIOR FILING DATE: 2001-11-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: BEDIAN, VAHE
APPLICANT: GLADUE, RONALD P.
APPLICANT: GLADUE, RONALD P.
APPLICANT: JIA, XIAO-CHI
APPLICANT: JIA, XIAO-CHI
APPLICANT: FENG, XIAO
TITLE OF INVENTION: ANTIBODIES TO CD40
FILE REFERENCE: ABX-PF/3 US
CURRENT APPLICATION NUMBER: US/11/211,917
CURRENT FILING DATE: 2005-08-25
PRIOR APPLICATION NUMBER: US/10/292,088
PRIOR FILING DATE: 2002-11-08
PRIOR FILING DATE: 2001-11-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Application US/1121191
No. US20060093600A1
                                                                                                                                                                                                                                                                                                                                                            15;
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     US20060093600A1
                                                                                                                                                                                                                                                                                                                                  th 56.6%; Similarity 70.1%; 82; Conservative 1
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Similarity 67.8%;
80; Conservative 1
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JEQ ID NOS: 14
JENGTH: 121
TYPE: PRT
JORGANISM: How
US-11-211-917-
Beer
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SOFTWARE: Patentin Ver. 2
SEQ ID NO 26
Publication No. US20060093
GENERAL INFORMATION:
APPLICANT: BEDIAN, VAHE
APPLICANT: GLADUE, RONAI
APPLICANT: CORVALAN, JOS
APPLICANT: JIA, XIAO-CH3
                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 1
SOFTWARE: Patentin Ver.
SEQ ID NO 109
LENGTH: 118
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Best Local S
Matches 82
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Pred. No. 6.5e-25
3; Mismatches 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 412; DB 7
Pred. No. 4.6e-2
1; Mismatches
US-11-211-917-98

US-11-211-917-98

Sequence 98, Application US/11211917

Publication No. US20060093600A1

GENERAL INFORMATION:

APPLICANT: BEDIAN, VAHE

APPLICANT: GLADUE, RONALD P.

APPLICANT: CORVALAN, JOSE

APPLICANT: JIA, XIAO-CHI

APPLICANT: JIA, XIAO-CHI

APPLICANT: FENG, XIAO

TITLE OF INVENTION: ANTIBODIES TO CD40

FILE REFERENCE: ABX-PF/3 US

CURRENT APPLICATION NUMBER: US/11/211,917

CURRENT FILING DATE: 2002-11-08

PRIOR FILING DATE: 2001-11-09

NUMBER OF SEQ ID NOS: 147

SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 98

LENGTH: 121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 66, Application US/11211917
; Publication No. US20060093600A1
; GENERAL INFORMATION:
; APPLICANT: BEDIAN, VAHE
; APPLICANT: GLADUE, RONALD P.
; APPLICANT: GLADUE, RONALD P.
; APPLICANT: JIA, XIAO-CHI
APPLICANT: JIA, XIAO-CHI
APPLICANT: FENG, XIAO
; TITLE OF INVENTION: ANTIBODIES TO CD40
; FILE REFERENCE: ABX-PF/3 US
; CURRENT APPLICATION NUMBER: US/11/211,917
; CURRENT FILING DATE: 2005-08-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US/10/292,088
PRIOR FILING DATE: 2002-11-08
PRIOR APPLICATION NUMBER: 60/348,980
PRIOR FILING DATE: 2001-11-09
NUMBER OF SEQ ID NOS: 147
SOFTWARE: Patentin Ver. 2.1
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illarity 68.9%;
Conservative 13
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82; Conservative
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Pred. No. 6.7e-25;
7; Mismatches 14
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1.1e-24;
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                                                                                                                        US-10-981-300-53
; Sequence 53, Application US/10981300
; Publication No. US20060093599A1
; GENERAL INFORMATION:
; APPLICANT: GIORGIO SENALDI
; APPLICANT: GADI GAZIT-BORNSTEIN
; TITLE OF INVENTION: ANTI-PROPERDIN ANTIBODIES
; TITLE OF INVENTION: FOR MAKING AND USING THE
; FILE REFERENCE: ABGX-005
; CURRENT APPLICATION NUMBER: US/10/981,300
; CURRENT FILING DATE: 2004-11-03
; NUMBER OF SEQ ID NOS: 71
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 53
; LENGTH: 115
; ORGANISM: Homo sapiens
US-10-981-300-53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 82, Application US/11211917
Sequence 82, Application US/11211917
GENERAL INFORMATION:
APPLICANT: BEDIAN, VAHE
APPLICANT: GLADUE, RONALD P.
APPLICANT: GLADUE, RONALD P.
APPLICANT: CORVALAN, JOSE
APPLICANT: JIA, XIAO-CHI
APPLICANT: JIA, XIAO-CHI
APPLICANT: JIA, XIAO-CHI
APPLICANT: JIA, XIAO-CHI
CURRENT FENG, XIAO
FILE REFERENCE: ABX-PF/3 US
CURRENT APPLICATION NUMBER: US/11/211,917
CURRENT APPLICATION NUMBER: US/10/292,088
PRIOR APPLICATION NUMBER: 60/348,980
PRIOR FILING DATE: 2001-11-09
PRIOR FILING DATE: 2001-11-09
NUMBER OF SEQ ID NOS: 147
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 82
LENGTH: 121
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ilarity 67.5%;
Conservative 1
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68.1%;
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US-11-211-917-82
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Matches 79; Conser
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US-11-211-917-82
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Matches 81
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updates/sec Search time 135.333 Seconds (without alignments) 391.900 Million cell upday

US-09-889-936A-8 606 1 SPCT score: Title: Perfect

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Scoring table:

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residues 457216429 2589679 segs, Searched:

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summaries

Database

A Geneseq 8:\*

1: geneseqp19808:\*

2: geneseqp20008:\*

4: geneseqp20018:\*

5: geneseqp20028:\*

6: geneseqp2003as:\*

7: geneseqp2003bs:\*

9: geneseqp20058:\*

10: geneseqp20058:\* 

results predicted by chance to have a to the score of the result being printed, of the total score distribution. greater than or equal to s derived by analysis of 햣 the number Ø % . score gand is Pred

## SUMMARIES

Description	1567	Add94117 Mouse HUI	7 Huma	Adz83525 AntiEpCam	60 Mous	83445 CD3 8	v66137 Anti-CD	dv66133 Anti-CD	dv66135 Anti-CD	dz8343	dv66116 Anti-CD	ab62864 Anti-S	8603 MCPC li	ar68742 MAb McP	58483 Murine	i01145 Murine	ar52036 Light ch	Ado32152 Mouse ant	6 Murine	02 FAB lig	<b>6</b> 8	S	Aar39336 scFv frag
ID	AAB15673	ADD94117	ADV66167	ADZ83525		ADZ83445	ADV66137	ADV66133	13	ADZ83436		AAB62864	3860	74	AAW58483		AAR52036	AD032152	AED64796	AAR53802		AAP80460	AAR39336
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541	AAR90834 AAR98479	4245	AEF80923 AAY06915	1379		ADG32320	AAB28395	AAU77947	ABO10744	ABO10756	ABR44700	ABR44688	ADQ90822	ADU67950	ADU67970	AED41851	AED41871	ADV66175
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## ALIGNMENTS

stric immunoassay; 2,4,6-trinitrotoluene; TNT. region. chain variable ; 5B3 antibody; IgG1; piezoelectric molecule explosive detection; 2,4,6 Z 116 2000WO-IL000048 99IL-00128212 standard; protein; entry) light antibody (first WO200043774-A2. 25-JAN-2000; 25-JAN-1999; 08-JAN-2001 27-JUL-2000 Murine 5B3 AAB15673; AAB15673 Mouse; small n 87 Mus RESULT 1 AAB15673 X

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2 Eshhar H Willner

2000-524259/47 B; AAA74605. N-PSDB; WPI

especially explosives comprises small molecules, Apparatus for detecting a piezoelectric crystal

90pp; English 3В; Disclosure; Fig The present sequence is the light chain variable region of 5B3 antibody. The cDNA encoding this sequence was obtained from total RNA extracted from a 5B3 hybridoma by RT-PCR. 5B3 is an IgG1 antibody derived from a TNP-KLH immunised mouse. Its binding to TNP or DNS antigen can be blocked by very low amounts of TNT and it can therefore be used in a method for detecting small assayed explosive molecules. Molecules are detected using a piezoelectric sensor. Piezoelectric immunoassaying in liquid phase allows stationary and flow analysis of an aqueous sample. The method is sufficiently sensitive for detection of low molecular weight molecules

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention relates to a novel grafted antibody or its functional fragment comprising one or more complementarity determining regions (CDRs) of a defined light CDR and a heavy CDR with at least one amino acid (aa) substitution where the antibody has specific binding activity for a cryptic collagen epitope. The growth of all solid tumours requires new blood vessel growth, angiogenesis, inhibition of which is an approach to limiting tumour growth. The invention may allow development of therapeutics with a cytostatic activity as a collagen agonist or antagonist. The invention is useful for diagnosing and treating disorders associated with angiogenesis, tumour growth and/or cancer metastasis. The present sequence is the partial amino acid sequence of the mouse anti-cryptic collagen site antibody HUIV26 variable region light chain used during the creation of the antibody of the invention.
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                                                                           SRGDIVMTQSPSSLSVSAGEKVTMSCKSSQSLLNSRNQKNYLAWYQQKPGQPPKLLIYGV
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                                                                                                                                                                                                                                                                                                                                                                          chain;
                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                complementarity
treating disorders
or cancer metastasi
                                                                                                                            cytostatic;
                                                                                                                                                                                                                                                                                                                      grafted antibody; complementarity determining region; CDR; ligh heavy CDR; cryptic collagen epitope; solid tumour; new blood vessel growth; angiogenesis; tumour growth; cytostaticollagen agonist; collagen antagonist; cancer metastasis; anti-cryptic collagen; antibody; HUIV26; variable region light
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                                                                                                                                                                                                                                                                                               acid
                         Length 116
                                                   Indels
                                                                                                                                                                                                                                                                                                amino
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antibody with one or more com
useful for diagnosing and tre
ogenesis, tumor growth and/or
                                                                                                                                                                                                                                                                                                chain partial
                                      .;
O 0
                         ; DB 3;
2.4e-43;
                                                 Mismatches
                        Score 606;
Pred. No. 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Broek
                                                                                                                                                                                                                                                                                               region light
                                                                                                                                                                                                                      standard; protein; 113
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New cryptic collagen antibody determining regions, useful for associated with angiogenesis,
                        100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2002WO-US038147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2001US-00995529
2001US-00011250
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                                                 Conservative
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                                                                                                                                                                                                                                                                      (first
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N-PSDB; ADD94116.
                                     Similarity
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                                                                                                                                                                                                                                                                                                                                                                                        murine
                                                                                                                                                                                                                                                                                                                                                                                                                 musculus.
  116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CELL
                                                                                                                                                                                                                                                                                               Mouse HUIV26
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06-DEC-2001;
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                                                 116;
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  Seguence
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                                                                                                                                                                                                                                                                                                                                                                                        mouse;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention comprises a composition that contains a bispecific single chain antibody consisting of at least two domains, where one of domains binds to human cell adhesion molecule (EpCAM) antigen, and the second domain binds to human CD3 antigen. The bispecific antibody construct of the invention is useful for the prevention, treatment or amelioration of a tumorous disease, such as an epithelial or minimal residual cancer. The present amino acid sequence represents an antibody light chain variable domain that is specific for human EpCAM.
                          PKLLIYGVFIR
                                    tumorous
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for preventing, treating or ameliorating
in epithelial or minimal residual cancer.
                          DIVMTQSPSSLSVSAGEKVTMSCKSSQSLLNSRNQKNYLAWYQQKPGQP
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         Indel
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Pred. No. 3.1e-38;
; Mismatches
Pred. No. 3.1e-38; Mismatches
                                                                                                                                                                                                               epithelial clight chain
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                                                                                                                                                                                                                                                                                                                                                                     Brischwein K,
U, Baeuerle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    227pp; English.
                                                                                                                                                                                            light
                                                                                                                                     Z
                                                                                                                                                                                                               in antibody;
cytostatic;
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         1;
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B, Lenkkeri-Schuetz
                                                                                                                                                                                             antibody
                                                                                                                                                                                                                                                                                                     2004WO-EP005687
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                89.9%;
92.0%;
Local Similarity 92.9%;
ses 105; Conservative
                                                                                                                                     standard; protein;
                                                                                                                                                                                                                                                                                                                      2003EP-0001213
2003EP-0001213
                                                                                                                                                                           entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    90;
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                                                                                                                                                                                                               fic single chatumor; cancer;
                                                                                                                                                                                             Epcam-specific
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 10; SEQ ID NO
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                                                                                                                                                                                                                                                                                                                                                   (MICR-) MICROMET AG
                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2005-021271/02
                                                                                                                                                                                                                                                                                                                                                                                                                              New pharmaceutical
construct, useful f
disease, such as ar
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104; Conser
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                                                                                                                                                                                                                                                                                                                      31-MAY-2003;
31-MAY-2003;
                                                                                                                                                                                                                                                                                                    26-MAY-2004;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 113
                                                                                                                                                                                                                                             Unidentified
                                                                                                                                                                                                                                                                                  09-DEC-2004
                                                                                                                                                                          24-FEB-2005
                                                                                                                                                                                                              bispecific
EpCAM; tumo
                                                                                                                                                                                                                                                                                                                                                                                Kohleisen
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                                                                                                                                    ADV66167
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                                                                                                                                                      ADV66167
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The invention relates to a cytotoxically active CD3 specific binding construct an Ig-derived second binding domain. The CD3 specific binding construct above or the construct produced by the process, nucleic acid molecule, vector, or host is useful for the preparation of a pharmaceutical composition for the prevention, treatment, or amelioration of a proliferative disease, a tumor, an inflammatory disease, an immunological disorder, an autoimmune disease, an inflammatory disease, viral disease, allergic reactions, parasitic reactions, graft-versus-host diseases, or host-versus-graft diseases. The cytotoxically active CD3 specific binding construct is useful for treating, preventing, or ameliorating proliferative disease, a tumor, an inflammatory disease, an immunological disorder, an autoimmune disease, an inflammatory disease, viral disease,
                                                                                                                                inflammation, immune disorder, infection, allergy;
sus host disease, Cytostatic, Antiinflammatory;
pressive, Virucide, Antibacterial; Antiallergic, Antiparasitic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   New cytotoxically active CD3 specific binding construct comprises a firs domain specifically binding to human CD3 and an Ig-derived second binding domain, useful for treating, preventing, or ameliorating, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   proliferative disease, a tumor, an inflammatory disease, an immunological disorder, an autoimmune disease, an inflammatory disease, an immunological allergic reactions, parasitic reactions, graft-versus-host diseases, or acid sequence of an anti-EpCam single chain antibody
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                                                                                                                                                                                                                                                                                                                                                                    Baeuerle
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Pred. No. 3.1e-38;
1; Mismatches 5;
                                                                                                                                                                                                                                                                                                                                                                    B, Lenkkeri-Schuetz U, Williams S;
                                                                                                        ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 147; 639pp; English.
                                                                                                        SEQ
                          Z
                         113
                                                                                                       antibody
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ilarity 92.0%;
Conservative
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                                                                                                                                                                                                                                                                                                              2003EP-00023581
                         standard; protein;
                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                      Kohleisen
                                                                                                         chain
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           disease
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B; ADZ83524.
                                                                                                                                             graft versus host d
Immunosuppressive;
EpCam; antibody.
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                                                                                                         single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 113 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            proliferative
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                                                                                                                                                                                                                                                                                   15-OCT-2004;
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8
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                                                                                                        AntiEpCam
                                                                                                                                    neoplasm;
                                                                                                                                                                                                    Synthetic
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Best Local S
Matches 104
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                        ADZ83525
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                                                   ADZ83525
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                                                                                                                                                                          EpCam;
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The invention relates to a method of identifying binding site domains (BSD) that retain the capacity of binding to a predetermined epitope when positioned C-terminal of at least one further domain in a recombinant bion positioned C-terminal of at least one further domain in a recombinant bion positioned C-terminal of a biological display system as part of a fusion protein for binding to a predetermined epitope, where the fusion protein comprises an additional domain positioned N-terminal of the BSD and an amino acid sequence that mediates anchoring of the fusion protein to the surface of the display system; and (b) identifying a BSD that binds to the predetermined epitope. The method is useful to identify bion or multivalent polypeptides that comprise antibody binding sites capable of efficiently binding to the corresponding antigen. The polypeptides or antibodies identified by the method are useful therapeutically and diagnostically, for e.g. cancer and autoimmune diseases. ScFv-antibody fragments that bind independently of their position within bifunctional single-chain fusion proteins can be isolated from combinatorial antibody libraries using the new in vitro method. Sequences AAY17957-965 represent
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                                                                                                                        cancer;
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                                                                                                                                                                                                                                                                                                                                                                           domains
                                                                                                                       therapeutic;
mouse.
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                                                                                                                       fusion protein; tsingle-chain Fv;
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Pred. No. 6.7e-38
; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                           identification
                                                                                                                                                                                                                                                                                                              Zett1
                                                                                                                       BSD; epitope;
scFv-antibody;
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                                                                                                                                                                                                                                                                                                                                                                                                                 6.6; 152pp; English
                       248
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ilarity 92.0%;
Conservative
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                                                                                                                                                                                                                                                                                                                Borschert
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AD283445
ID AD283445 standard; protein;
                       protein;
                                                                                                                                                                                                                                                                                                                                                                                           epitope
                                                                                                                        domain; BSD;
                                                                                                                                                                                                                                                                                                                                                                          Phage display system for capacity to bind an epito
                                                                                                4-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       scFV fragments
                                                                                                                                                                                                                                                                                                                                      WPI; 1999-338004/28
N-PSDB; AAX77243.
                                                                      (first
                                                                                                fragment
                                                                                                                                  disease;
                      standard;
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                                                                                                                        site
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                                                                                                                                                                                                                                                            17-NOV-1997;
                                                                      04-AUG-1999
                                                                                                                                                                                                            27-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              104;
                                                                                              SCFV
                                                                                                                                    autoimmune
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Best Local S
Matches 104
                                              AAY17960;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
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                       AAY17960
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                                                                                                                                                                                    WO992581
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                                                                                                                       Binding
                                                                                                Mouse
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RESULT 5
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thain antibody; epithelial cancer; cytostatic.

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antibody

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The invention comprises a composition that contains a bisp chain antibody consisting of at least two domains, where o binds to human epithelial cell adhesion molecule (EpCAM) a second domain binds to human CD3 antigen. The bispecific a construct of the invention is useful for the prevention, tamelioration of a tumorous disease, such as an epithelial residual cancer. The present amino acid sequence represent single chain antibody of the invention.
                                                                                                                                                                                                                                                                                                                                                  pharmaceutical composition having a bis
struct, useful for preventing, treating
sase, such as an epithelial or minimal r
                           Anti-CD3-anti-EpCAM
                                                                                                                                                                                                                                                                                                            WPI; 2005-021271/02
N-PSDB; ADV66136.
                                                    bispecific single EpCAM; CD3; tumor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 500 AA;
                                                                                                                      WO2004106383-A1
                                                                                                                                                                                                  31-MAY-2003;
31-MAY-2003;
                                                                                                                                                                          26-MAY-2004;
                                                                                            Unidentified
 24-FEB-2005
                                                                                                                                                                                                                                                                                                                                                                 construct,
                                                                                                                                                                                                                                                                                    Kohleisen
                                                                                                                                                                                                                                                                                                                                                                                                         Claim 12;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a cytotoxically active CD3 specific binding construct comprises a first domain specifically binding to human CD3 and and generated second binding domain. The CD3 specific binding construct above or the construct produced by the process, nucleic acid molecule, vector, or host is useful for the preparation of a pharmaceutical composition for the prevention, treatment, or amelioration of a proliferative disease, a tumor, an inflammatory disease, an immunological disorder, an autoimmune disease, an infectious disease, viral disease, allergic reactions, parasitic reactions, graft-versus-host diseases, or host-versus-graft disease. The cytotoxically active CD3 specific binding construct is useful for treating, preventing, or ameliorating proliferative disease, a tumor, an inflammatory disease, an immunological disorder, an autoimmune disease, an inflammatory disease, viral disease, allergic reactions, parasitic reactions, graft-versus-host diseases, or host-versus-graft diseases. The present sequence represents the amino caid sequence of a CD3 specific binding construct.
                                                                                                                                                                                                                                                                                                                                                                             a first
binding
                                                                            nune disorder; infection; allergy;
Cytostatic; Antiinflammatory;
s; Antibacterial; Antiallergic; Antiparasitic
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                                                                                                                                                                                                                                                                                                Baeuerle
                                                                                                                                                                                                                                                                                                                                                                             ig construct comprises I an Ig-derived second ameliorating, e.g.
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1.3e-37;
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Williams S;
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                                                                               immune
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                                                                            neoplasm; inflammation; immusgraft versus host disease; CyImmunosuppressive; Virucide;
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                                                                                                                                                                                                                 2004WO-EP011646
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                           entry)
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                                                                                                                                                                                                                                                                                                          Hamilton AA,
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                                                    binding
                          (first
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N-PSDB; ADZ83444.
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                                                    specific
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                                                                                                                                                                                                                                                                                               Hofmeister
                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 20;
                                                                                                                                  Synthetic
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one of dom antigen, a antibody treatment

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227pp; English

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SEQ ID NO

chain antibody a tumorous

bispecific single cang or ameliorating the sidual cancer.

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Wolf

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Brischwein K, U, Baeuerle

y M, Offner S, Lenkkeri-Schuetz

Berry

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2004WO-EP005687

2003EP-000121 2003EP-000121

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1.4e-37;
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ore 545; DB ed. No. 1.4e Mismatches
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Pred.
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89.9%;
ilarity 92.0%;
Conservative
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       l Similarity
104; Conser
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EpCAM; CD3; tumor
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ed. No. 1.3 Mismatches

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89.9%; ilarity 92.0%; Conservative

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4 DIVMTQSPSSLSVSAGEKVTMSCKSSQSLLNSRNQKNYLAWYQQKPGQPPKLLIYGVFIR

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ADV66137 standard; protein; 500 AA

ADV66137

RESULT 7
ADV66137
ID ADV6
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WPI; 2005-021271/02
N-PSDB; ADV66134.
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graft vers
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                                                                                                                                                                         chain antibody consisting of at least two domains, where one of dom binds to human epithelial cell adhesion molecule (EpCAM) antigen, second domain binds to human CD3 antigen. The bispecific antibody construct of the invention is useful for the prevention, treatment amelioration of a tumorous disease, such as an epithelial or minima residual cancer. The present amino acid sequence represents a bispesingle chain antibody of the invention.
                                                                                                                                                                                                                                                                                                                                        497
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U, Baeuerle
                                                                              Baeuerle
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Pred. No. 1.4e
4; Mismatches
                                                                                                                                                                    comprises a composition that
                                                                   Brischwein
U, Baenerl
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B, Lenkkeri-Schuetz
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Lenkkeri-Schuetz
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2003EP-00012134
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2003EP-00012134
                2004WO-EP005687
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104; Conservative
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disease, such as an
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EpCAM; CD3;
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31-MAY-2003
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nain antibody
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versus host disease; Cytostatic; Antiinflammatory;
ssuppressive; Virucide; Antibacterial; Antiallergic; Antiparasiti
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minimal
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                                                                                               The invention comprises a composition that contains a bispecific sichain antibody consisting of at least two domains, where one of dom binds to human epithelial cell adhesion molecule (EpCAM) antigen, a second domain binds to human CD3 antigen. The bispecific antibody construct of the invention is useful for the prevention, treatment amelioration of a tumorous disease, such as an epithelial or minima residual cancer. The present amino acid sequence represents a bispesingle chain antibody of the invention.
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    chain
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 New pharmaceutical composition having a bispecific single c
construct, useful for preventing, treating or ameliorating
disease, such as an epithelial or minimal residual cancer.
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ilarity 92.0%;
Conservative
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The invention relates to a cytotoxically active CD3 specific binding construct comprises a first domain specifically binding to human CD3 and an Ig-derived second binding domain. The CD3 specific binding construct above or the construct produced by the process, nucleic acid molecule, vector, or host is useful for the preparation of a pharmaceutical composition for the prevention, treatment, or amelioration of a proliferative disease, a tumor, an inflammatory disease, an immunological disorder, an autoimmune disease, an infectious disease, viral disease, allergic reactions, parasitic reactions, graft-versus-host diseases, or host-versus-graft diseases. The cytotoxically active CD3 specific binding construct is useful for treating, preventing, or ameliorating
                                                                                                                                                                                                                                                                                                                                                      disease, an immunologica
lisease, viral disease,
                                                                                                                                                                                                                                                                                                                          construct is useful for treating, preventing, or ameliorating proliferative disease, a tumor, an inflammatory disease, an immunologidisorder, an autoimmune disease, an infectious disease, viral disease, allergic reactions, parasitic reactions, graft-versus-host diseases, host-versus-graft diseases. The present sequence represents the amino acid sequence of deimmunized construct 4-lxanti-CD3.
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Score 545; DB 9; Le
Pred. No. 1.4e-37;
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     89.9%;
92.0%;
               Conservative
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     Query Match
Best Local
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### ADV66116 standard; protein; 521 AA ADV66116; RESULT 11 ADV661

entry) (first 24-FEB-2005

SEQ ID antibody chain single Anti-CD3-anti-EpCAM bispecific

39

chain antibody; epithelial cell adhesion molecule; cancer; cytostatic. tumor; single bispecific E EpCAM; CD3;

Unidentified

WO2004106383-A1

09-DEC-2004

2004WO-EP005687 26-MAY-2004; 2003EP-00012133 -MAY-2003; -MAY-2003;

(MICR-) MICROMET AG

Raum Ą Wolf <u>ب</u> Brischwein K, U, Baeuerle y M, Offner S, Lenkkeri-Schuetz Berry M. Kohleisen B, Kufer

WPI; 2005-021271/02 N-PSDB; ADV66115.

antibody pharmaceutical composition having a bispecific single chain antibortruct, useful for preventing, treating or ameliorating a tumorous sase, such as an epithelial or minimal residual cancer. construct, disease, Ze Z

English. Claim 12; SEQ ID NO 39; 227pp;

composition that contains a bispecific single of at least two domains, where one of domains cell adhesion molecule (EpCAM) antigen, and the The invention comprises a chain antibody consisting binds to human epithelial

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This invention relates to sialoadhesin factor-1 (SAF-1) monoclonal antibodies 13G3 and 11G4. A neutrophil-priming dose of SAF-1 agonist antibodies 13G3 and 11G4. A neutrophil-priming dose of SAF-1 agonist cantibody or a pharmaceutical composition comprising the antibody is useful for treating or preventing an infectious disease state such as costeomyelitis, deep seated wound infections, sepsis, antibiotic resistant bathogens and parasites such as Listeria monocytogenes, Salmonella and cathogens and parasites such as Listeria monocytogenes, Salmonella and Leishmania in a mammal. Nucleic acid sequences encoding the variable introduction of specific changes within the nucleic acid sequences introduction of the resulting modified or fusion regions, and for incorporation of the resulting modified or fusion capins acid sequence into a plasmid for expression. The CDR-encoding cregions comprising silent mutations are used in construction of humanized
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ion of humanized
                                      a bispecific
                                                                                                                                                         63
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                                                                                                                                                                                                                                                                                                                                                                                                        13G3; 11G4
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                                                                                                                                                                                                                                                                                                                                                                              region protein
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             treatment or or or minimal
                                                                                                                                                         YGVFIR
                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          factor-1 agonist antibody having the characteristic ody 13G3 or 11G4 for treating osteomyelitis, deep sevand antibiotic resistant bacterial infections.
                                                                                                                                                                                                            116
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                                                                                                                                                        DIVMTOSPSSLSVSAGEKVTMSCKSSQSLLNSRNOKNYLAWYQQKPGQPPKLLI
                                                                                                                                                                                                                                                                                                                                                                                                      nal antibody; 1 monocytogene; ion; mouse.
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0
                                      represents
                                                                                                     Length 521
             the prevention, tas an epithelial
  bispecific
                                                                                                                                                                                                                                                                                                                                                                             variable
                                                                                                                                Indels
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                                                                                                                                                                                                                                                                                                                                                                                                     Sialoadhesin factor-1; SAF-1; anti-SAF-1 monoclonal e osteomyelitis; wound infection; sepsis; Listeria mono Salmonella; Leishmania; kappa chain variable region;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Taylor
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ر.
                                       sequence
                                                                                                                                                                                                                                                                                                                                                                              antibody 13G3 kappa chain
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  The
                                                                                                      DB 9,
                                                                                                      ore 545; DB
ed. No. 1.4e
Mismatches
second domain binds to human CD3 antigen. construct of the invention is useful for tamelioration of a tumorous disease, such residual cancer. The present amino acid staingle chain antibody of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             King AG,
                                                                                                      Score
Pred.
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                                                                                                                                                                                                                                                                                                  standard; protein; 114
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                                                                                                    89.9%;
ilarity 92.0%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                       entry)
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monoclonal antibody 13G3 (
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                                                                                                                                                                                                                                                                                                                                                                               monoclonal
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N-PSDB; AAF62143.
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Rest Local Similarity
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                                                                              521
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                                                                                                                            104;
                                                                                                                                                                                                                                                                                                                                                                              Anti-SAF-1
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                                                                              Sequence
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                                                                                                   63
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                                                                                                            DIVMTOSPSSLSVSAGEKVTMSCKSSOSLLNSRNOKNYLAWYQQKPGQPPKLLIYGVFIR
                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           retain
       present sequence
kappa chain variable
                                                                                                                                                                                                                                                                                               affinity;
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                                                                                                                                                           113
                                                                                                                                                                                                                                                                                                                                                                 of hypervariable loop
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Уq
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cof auto-immune diseases reduce immunogenicity but
                                                            Length 114
                                                                                                                                                                                                                                                                                               Antibody; variable domain; light; L; heavy; H; consensus; antigen; immunogenicity; humanisation; framework.
                                                                                Indels
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                                                                               9
                                                            ore 540; DB 4;
ed. No. 8.2e-38;
Mismatches 6;
    antibodies or other engineered antibodies. The represents anti-SAF-1 monoclonal antibody 13G3 region amino acid sequence
                                                                                                                                                                                                                                                                                                                                                                "variation in length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  pn. used for treatment critical residues to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 85; 160pp; English
                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                             Score
Pred.
                                                                                                                                                                                                          standard; protein; 113
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                                                           89.1%;
ilarity 91.2%;
Conservative
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/note= "
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(first en
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                                                                       l Similarity
103; Conser
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replacement of cri
binding affinity,
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                                            Sequence 114 AA;
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Misc-difference
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                                                                                                                                                                                                                                                25-MAR-2003
28-OCT-1993
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                                                                                                                                                                                                                                                                           MCPC light
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                                                             Query Match
Best Local S
Matches 103
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possible interchain disulfide bond"
                                                       DIVMTQSPSSLSVSAGEKVTMSCKSSQSLLNSRNQKNYLAWYQQKPGQPPKLLIYGVFIR
                                      Gaps
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| label= CDR3
| note= "complementarity determining region
| note= "complementarity determining region
                                                                                                                                                                                                                                                                                                                                                                                    "complementarity determining region
                                                                                                                                                                                                                                                              disulfide bond; heavy chain; engineering; ligand binding m
                   113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               region
                                     Indels
                  Length
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possible interchain disulfide bond"
56. .62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "complementarity determining
                           ore 536; DB 2; ed. No. 1.8e-37; Mismatches 7
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Pred.
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1. .23
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                                                                                                                                                                                                                                                            Monoclonal antibody; McPC603; variable region; Fv; antibody immunotoxin.
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|abel= CDR2
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                                                                                                                                                                     AAR68742 standard; protein;
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63. .94
/label= FR3
                 ch 88.4%;
il Similarity 91.2%;
103; Conservative
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(first entry)
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/label= 1
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/label= (
/note= "c
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/label= 1
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24. .40
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113 AA;
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15-JUL-1995
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                 Query Match
Best Local S
Matches 103
Sequence
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                                                                                                                                                                                                                                    9
                                      having a disulphide bond linking of variable regions of ligand binding
                                                                                                                                                                                                                 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antibodies with
                                                                                                  The polypeptide given in AAR68742 corresponds to the light chain of MAD McPC603. This MAD was used to generate a model of MAD B3(Fv), which has been used in the construction of disulfide bond-stabilized Fv fragments used as immunotoxins. (Updated on 25-MAR-2003 to correct PN field.)
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                                                                                                                                                                                                                                                                                                                                                                                                            mouse; CD5; anti-CD5 antibody; immunoglobulin;
cic; immunoconjugate; fusion protein; psoriasis
e; rheumatoid arthritis; type I diabetes.
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Pred. No. 1.8e-37
3; Mismatches
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92WO-US010906.
93US-00082842.
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ilarity 91.2%;
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cytotoxic;
disease; rh
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                                                                                                                                                                                                                                                                                                                                                                                          antibody
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                   1995-036408/05
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103; Conserv
                                    ligand-binding
                                                                                                                                                                                                                                                                                                                              standard;
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CD)
                                                  framework regions
                                                                               Disclosure; Page
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                                                                                                                                                    Sequence 113
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14-DEC-1992;
23-JUN-1993;
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                                                            moieties.
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Best Loc
Matches
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AAW58483
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AAW58480), and (b) a modified heavy chain variable region (see AAW58479 or AAW58481), where AAW58478 and AAW58479 are humanised forms of the H65 light and heavy chain variable domains with low risk amino acid substitutions [i.e. low risk of reducing antigen-binding specificity.] and AAW58480 and AAW58481 are humanised forms of the H65 light and heavy chain variable domains with moderate risk amino acid substitutions and are present in humanised H65 antibody he3 (ATCC HB 11206). The method is useful for treating autoimmune diseases, especially systemic lupus erythematosus, rheumatoid arthritis, psoriasis or type I diabetes. The present sequence represents the murine MCPC antibody light chain variable
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Pred. No. 1.8e-37
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Local Similarity 91.2%;
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Search

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version - 2006
GenCore (c) 1993
       Copyright
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using sw model protein search, OM protein

17:40:37 2006, 19, Мау Run on:

Search time 22.0952 Seconds (without alignments) 505.138 Million cell updates/sec

.....CQNDHIYPYTFGGGTKLEIK SRGDIVMTQSPSSLSVSAGE.. US-09-889-936A-8 606 Title: Perfect score: Sequence:

116

BLOSUM62 Gapop 10.0 , Scoring table

Gapext

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residues 96216763 283416 seqs, Searched

283416 hits satisfying chosen parameters: of Total number

Minimum DB Maximum DB

length: 0 length: 2000000000 seq

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

PIR Database

R\_80:\* pir1:\* pir2:\* pir3:\* 1 0 E 4

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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# ALIGNMENTS

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Similarity 92.5%;
99; Conservative
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C;Species: Mus musculus (house mouse)
C;Date: 16-Sep-1992 #sequence_revision 16-Sep-1992 #text_change 21-J
C;Accession: PL0264
R;Shlomchik, M.; Mascelli, M.; Shan, H.; Radic, M.Z.; Pisetsky, D.; J.
J. Exp. Med. 171, 265-297, 1990
A;Title: Anti-DNA antibodies from autoimmune mice arise by clonal exp A;Reference number: PL0231; MUID:90111618; PMID:2104919
A;Accession: PL0264
A;Molecule type: mRNA
A;Residues: 1-113 <SHL>
A;Cross-references: UNIPARC: UPI0000176AFB
C;Superfamily: immunoglobulin V region; immunoglobulin
F;1-23/Region: framework 1
F;16-96/Domain: immunoglobulin homology <IMM>F;24-40/Region: complementarity-determining 1
F;6-62/Region: complementarity-determining 7
F;56-62/Region: framework 3
F;56-103/Region: framework 3
F;56-103/Region: framework 3
F;56-103/Region: framework 3
F;104-11:**
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Cross-references: UNIPARC:UPI0000176CD4
Superfamily: immunoglobulin V region; immunoglobulin homology
Keywords: heterotetramer; immunoglobulin
16-96/Domain: immunoglobulin homology <IMM>
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C; Species: Mus musculus (house mouse)
C; Date: 10-Feb-1989 #sequence_revision 10-Feb-1989 #text_char
C; Accession: E30538
R; Clafiln, J.L.; Berry, J.
J. Immunol. 141, 4012-4019, 1988
A; Title: Genetics of the phosphocholine-specific antibody res
A; Reference number: A30534; MUID:89035545; PMID:3141511
A; Accession: E30538
A; Status: preliminary; nucleic acid sequence not shown; not c
A; Molecule type: mRNA
A; Residues: 1-112 < CLA>
A; Cross-references: UNIPARC:UPI0000176CD4
C; Superfamily: immunoglobulin V region; immunoglobulin homolox
C; Keywords: heterotetramer; immunoglobulin:
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Pred. No. 2.4e-40;
2; Mismatches 7;
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6.8e-40;
Score 536; DB 2;
Pred. No. 2.4e-40;
; Mismatches
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Pred. No. 6.8e
4; Mismatches
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ilarity 90.3%;
Conservative
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92.0%;
Match
Local Similarity 92.0%;
nes 103; Conservative
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G30535
Ig kappa chain V region (2G6) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 23-Feb-1989 #sequence_revision 23-Feb-1989 #text_change 21-Jar
C;Accession: G30535
R;Claflin, J.L.; Berry, J.; Flaherty, D.; Dunnick, W.
J. Immunol. 138, 3060-3068, 1987
A;Title: Somatic evolution of diversity among anti-phosphocholine antiA; Reference number: A30556; MUID:87196439; PMID:3106498
A;Accession: G30535
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-107 < CLA>
A;Coss-references: UNIPARC:UPI0000176AF6
A;Note: the sequence was determined from the differentiated gene
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;16-96/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                     Ig kappa chain V region (6D10) - mouse (fragment)
C; Species: Mus musculus (house mouse)
C; Date: 23-Feb-1989 #sequence_revision 23-Feb-1989 #text_change 21-Ja:
C; Date: 23-Feb-1989 #sequence_revision 23-Feb-1989 #text_change 21-Ja:
C; Accession: E30535
R; Claflin, J.L.; Berry, J.; Flaherty, D.; Dunnick, W.
J. Immunol. 138, 3060-3068, 1987
A; Title: Somatic evolution of diversity among anti-phosphocholine ant.
A; Reference number: A30556; MUID:87196439; PMID:3106498
A; Accession: E30535
A; Accession: E30535
A; Residues: 1-108 < CLA>
A; Residues: 1-108 < CLA>
A; Residues: 1-108 < CLA>
A; Residues: UNIPARC:UPI0000176AED
A; Cross-references: UNIPARC:UPI000176AED
A; Note: the sequence was determined from the differentiated gene
C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
F; 16-96/Domain: immunoglobulin homology < IMM>
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Pred. No. 1.6e-38
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6e-39;
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1; Mismatches
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                                                                      C; Species: Mus musculus (house mouse)
C; Species: Mus musculus (house mouse)
C; Accession: D30535
R; Claflin, J.L.; Berry, J.; Flaherty, D.; Dunnick, W.
J. Immunol. 138, 3060-3068, 1987
A; Title: Somatic evolution of diversity among anti-phosphocholine ant A; Reference number: A30556; MUID:87196439; PMID:3106498
A; Accession: D30535
A; Accession: D30535
A; Molecule type: DNA
A; Molecule type: DNA
A; Residues: 1-107 < CLA>
A; Residues: 1-107 < CLA>
CLA>
C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Superfamily: immunoglobulin
C; Keywords: heterotetramer; immunoglobulin
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C; Species: Muss wusculus (house mouse)
C; Species: Muss wusculus (house mouse)
C; Species: Muss wusculus (house mouse)
C; Accession: B30535
R; Claflin, J.L.; Berry, J.; Flaherty, D.; Dunnick, W.
J. Immunol. 138, 3060-3068, 1987
A; Title: Somatic evolution of diversity among anti-phosphocholine ant A; Reference number: A30556; MUID:87196439; PMID:3106498
A; Reference number: A30556; MUID:87196439; PMID:3106498
A; Accession: B30535
A; Molecule type: DNA
A; Residues: 1-107 < CLA>
A; Cross-references: UNIPARC:UPI0000176AF5
A; Note: the sequence was determined from the differentiated gene C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
F; 16-96/Domain: immunoglobulin homology < IMM>
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Pred. No. 6.7e-38;
; Mismatches 8;
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Pred. No. 6.7e-38;
Mismatches
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(house mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              - mouse
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Similarity 90.7%;
97; Conservative
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98; Conserv
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Best Local S
Matches 98
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Best Local S
Matches 97
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C; Date: 16-Sep-1992 #sequence_revision 16-Sep-1992 #text_change 21-Jan-2000
C; Accession: PLO263
R; Shlomchik, M: Mascelli, M: Shan, H:; Radic, M.Z.; Pisetsky, D.; Marshak-Roth J. Exp. Med. 171, 265-297, 1990
A; Title: Anti-DNA antibodies from autoimmune mice arise by clonal expansion and A; Reference number: PLO231; MUID: 90111618; PMID: 2104919
A; Reference number: PLO231; MUID: 90111618; PMID: 2104919
A; Resion: PLO263
A; Molecule type: mRNA
A; Residues: 1-13 <81L.>
A; Resion: framework unmunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
F; 16-96 Domain: framework 1
F; 16-96 Domain: framework 2
F; 56-94 Region: complementarity-determining 1
F; 16-95 Region: complementarity-determining 2
F; 56-94 Region: complementarity-determining 2
F; 56-94 Region: complementarity-determining 2
F; 56-94 Region: complementarity-determining 2
F; 56-94 Region: complementarity-determining 2
F; 56-13 Region: complementarity-determining 2
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F; 56-13 Region: complementarity-determining 2
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                                                                                                                                          21-Jan-
                                                                                                          C; Species: Mus musculus (house mouse)
C; Species: Mus musculus (house mouse)
C; Date: 23-Feb-1989 #sequence_revision 23-Feb-1989 #text_change 21-Ja
C; Accession: F30535
R; Claflin, J.L.; Berry, J.; Flaherty, D.; Dunnick, W.
J. Immunol. 138, 3060-3068, 1987
A; Title: Somatic evolution of diversity among anti-phosphocholine ant A; Reference number: A30556; MUID:87196439; PMID:3106498
A; Reference number: A30556; MUID:87196439; PMID:3106498
A; Residues: Preliminary
A; Molecule type: DNA
A; Residues: 1-107 < CLA>
A; Cross-references: UNIPARC:UPI0000176AFA
A; Cross-references: UNIPARC:UPI000176AFA
C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
F; 16-96/Domain: immunoglobulin homology < IMM>
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Pred. No. 5.8e-38;
; Mismatches 10;
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2; Mismatches
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Similarity 86.7%;
98; Conservative
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ilarity 82.3%;
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Similarity 84.1%;
95; Conservative
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Local Similarity 83.2%;
nes 94; Conservative
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93; Conser
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Best Local S
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PMID:7498516
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C; Species: Mus musculus (house mouse)
C; Species: Mus musculus (house mouse)
C; Date: 31-Mar-1990 #sequence_revision 31-Mar-1990 #text_change 21-J.
C; Accession: A31790
R; Schulze-Gahmen, U.; Rini, J.M.; Arevalo, J.; Stura, E.A.; Kenten, J. Biol. Chem. 263, 17100-17105, 1988
A; Title: Preliminary crystallographic data, primary sequence, and bi. A; Reference number: A92686; MUID:89034213; PMID:3182835
A; Accession: A31790
A; Molecule type: mRNA
A; Residues: 1-220 <SCH>
A; Cross-references: UNIPARC:UPI0000114E12; GB:M23626; GB:J04061; NID C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
F; 16-96/Domain: immunoglobulin homology <IMM>
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C; Species: Mus musculus (house mouse)
C; Date: 29-Jul-1997 #sequence_revision 29-Aug-1997 #text_change 11-C; Accession: $68212
R; Takagi, M.; Kohda, K.; Hamuro, T.; Harada, A.; Yamaguchi, H.; Kan FEBS Lett. 375, 273-276, 1995
A; Title: Thermostable peroxidase activity with a recombinant antibo A; Reference number: $68211; MUID:96085223; PMID:7498516
A; Reference number: $68212
A; Katus: preliminary; nucleic acid sequence not shown
A; Molecule type: mRNA
A; Residues: 1-214 < TAK>
A; Cross-references: UNIPARC:UPI000017697E; EMBL:D29668
C; Superfamily: immunoglobulin V region; immunoglobulin homology
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Immunol. 138, 3060-3068, 1987
Title: Somatic evolution of diversity among anti-phosphocholine Reference number: A30556; MUID:87196439; PMID:3106498
Accession: C30535
Status: preliminary
                                                                                                                                                                                                                            gene
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(Cross-references: UNIPARC: UPI0000176AF4)
(Note: the sequence was determined from the differentiated general sequence was determined from the differentiated general sequence was determined from the differentiated general sequence was determined from the differentiated general sequence was determined from the differentiated general sequence was determined from the differentiated general sequence was determined from the differentiated general sequence was determined from the differentiated general sequence was determined from the differentiated general sequence was determined from the differentiated general sequence was determined from the differentiated general sequence was determined from the differentiated general sequence was determined from the differentiated general sequence was determined from the differentiated general sequence was determined from the differentiated general sequence was determined from the differentiated general sequence was determined from the differentiated general sequence was determined from the differentiated general sequence was determined from the differentiated general sequence was determined from the differentiated general sequence was determined from the differentiated general sequence was determined from the differentiated general sequence was determined from the differentiated general sequence was determined from the differentiated general sequence was determined from the differentiated general sequence was determined from the differentiated general sequence was determined from the differentiated general sequence was determined from the differentiated general sequence was determined from the differentiated general sequence was determined and determined general sequence was determined from the differentiated general sequence was determined and determined general sequence was determined and determined general sequence was determined general sequence was determined and determined general sequence was determined general sequence was determined general seque
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12;
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1.8e-37;
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Pred. No. 1.9e-36
6; Mismatches 13
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  J. Immunol. 138, 30 A; Title: Somatic ev A; Reference number: A; Accession: C30535 A; Status: prelimina A; Molecule type: DN A; Residues: 1-105 < A; Cross-references: A; Note: the sequenc C; Superfamily: immu C; Keywords: heterot F; 16-96/Domain: imm
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PT0407

Ig light chain V region (S107/VH11 group 1-4) - mouse (fragment)
C; Species: Mus musculus (house mouse)
C; Species: Mus musculus (house mouse)
C; Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 21-Jan C; Accession: PT0407
R; Behar, S.M.; Lustgarten, D.L.; Corbet, S.; Scharff, M.D.
J. Exp. Med. 173, 731-741, 1991
A; Title: Characterization of somatically mutated S107 VH11-encoded ant. A; Resion: PT0407
A; Molecule type: DNA
A; Residues: 1-113 < BEH>
A; Cross-references: UNIPARC: UPI0000176A05
C; Superfamily: immunoglobulin V region; immunoglobulin
F; 16-96/Domain: immunoglobulin homology
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Cross-references: UNIPARC:UPI0000116626; EMBL:Z31353; NID:g467!
Cross-references: UNIPARC:UPI0000116626; EMBL:Z31353; NID:g467!
Superfamily: immunoglobulin V region; immunoglobulin
Keywords: heterotetramer; immunoglobulin
16-96/Domain: immunoglobulin homology <IMM>
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S43103
Ig kappa chain V-J region (4B1 VL) - mouse (fragment)
C; Species: Mus musculus (house mouse)
C; Date: 06-Feb-1995 #sequence_revision 06-Feb-1995 #text
C; Accession: S43103
R; Gilbert, D.; Brard, F.; Margaritte, C.; Delpech, A.; T
submitted to the EMBL Data Library, March 1994
A; Description: An idiotype D23-bearing polyspecific, mux
A; Reference number: S42484
A; Accession: S43103
A; Residues: 1-112 <GIL>
A; Residues: 1-112 <GIL>
A; Cross-references: UNIPARC: UPI0000116626; EMBL: Z31353;
C; Superfamily: immunoglobulin V region; immunoglobulin h
C; Keywords: heterotetramer; immunoglobulin h
F; 16-96/Domain: immunoglobulin homology <IMM>
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ed. No. 7.6e-35;
Mismatches 11;
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1.4e-35;
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  485;
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### November 2005

Published\_Applications Nucleic Acid and Published\_Applications Amino Acid database searches now generate two sets of results each. The Published\_Applications databases have been split into two parts to reduce the amount of time required for their daily updates. This results in more machine time being available for processing searches.

Newly published applications will appear in the Published\_Applications\_New databases; older published applications make up the Published\_Applications\_Main databases.

Searches run against Nucleic Acid Published\_Applications produce two sets of results, with the extensions <code>.rnpbm</code> (Published\_Applications\_NA\_Nain) and <code>.rnpbn</code> (Published\_Applications\_NA\_New).

Searches run against Amino Acid Published\_Applications produce two sets of results, with the extensions <code>.rapbm</code> (Published\_Applications\_AA\_New).



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### **ALIGNMENTS**

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Distributed under the Creative Commons Attribution-NoDerivs License
                               05-JUL-2004, integrated into UniProtKB/TrEMBL.
05-JUL-2004, sequence version 1.
07-FEB-2006, entry version 11.
ScFv B8E5 protein (Fragment).
Name=scFv B8E5;
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Buarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Muridae; Mus.
NCBI TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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STRAINDER Balb/C;
PubMed=15485827; DOI=10.1074/jbc.M407213200;
Peter J.C., Wallukat G., Tugler J., Maurice D., Roegel Briand J.P., Hoebeke J.;
"Modulation of the M2 muscarinic acetylcholine receptor monoclonal anti-M2 receptor antibody fragments.";
J. Biol. Chem. 279:55697-55706(2004).
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length
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No. 2.4e-48
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                     255
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                    PRT;
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SMART; SM00406; IGv; 2.
PROSITE; PS50835; IG_LIKE;
Immunoglobulin domain.
NON TER
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Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

A Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

A Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

Mhiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,

Butterfield X.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,

"Generation and initial analysis of more than 15,000 full-length human
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Distributed under the Creative Commons Attribution-NoDerivs License
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi
Muroidea; Muridae; Mus.
NCBI_TaxID=10090;
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                                                                                                                                    WAP-TGF alpha model.
                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE. FVB/N; TISSUE=Mammary tumor. WAP-TGF alpha model
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Submitted (APR-2005) to the EMBL/GenBank/DDBJ databases
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                                                                                                                                                                                                                                                                                                                                                                99:16899-16903(2002)
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Pred. No. 1.3e-41;
0; Mismatches 14;
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                     into UniProtKB/TrEMBL
   240 AA
                                                                                                                           NUCLEOTIDE SEQUENCE.
STRAIN=Mix FVB/N; TISSUE=Mammary tumor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAH94049.1; -; mRNA
   PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; 2.
UNKNOWN
Q52L64 MOUSE PRELIMINARY; PRT; Q52L64; 24-MAY-2005, integrated into UniPr 24-MAY-2005, sequence version 1. 07-FEB-2006, entry version 10. Hypothetical protein. Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10;
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0; 19-1ike.
7; 19-ci.
16; 19-MHC.
36; 19-v.
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id. Sci. U.S.A.
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PROSITE; PS00290; IG_MHC;
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90; Conservative
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InterPro; IPR003599; Ig.
InterPro; IPR003110; Ig-
InterPro; IPR003597; Ig_
InterPro; IPR003596; Ig_
InterPro; IPR003596; Ig_
Pfam; PF07654; C1-Bet; I
SMART; SM00409; IG; 1.
SMART; SM00406; IG; 1.
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240 AA; 2
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STRAIN=Mix
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Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                          Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                        a cloned cDNA
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                                                                                                                                            15;
                                                                                                                               th Similarity 77.0%; Pred. No. 1e-40; 87; Conservative 11; Mismatches 15.
                                                                                                                   0647F1D17F236485
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NUCLEOTIDE SEQUENCE [MRNA].
MEDLINE=86041854; PubMed=2997713;
Marsh P., Mills F., Gould H.;
"Detection of a unique human V kappa IV ge probe.";
Nucleic Acids Res. 13:6531-6544(1985).
[2]
                                                                                                                                                                                                                                                                                                                                                                                                         EMBL; X02990; CAA26733.1; -; mRNA.
HSSP; P01625; 1LVE.
SMR; P06314; 21-134.
LinkHub; P06314; -.
GO; GO:0005576; C:extracellular region; NA GO; GO:0003823; F:antigen binding; NAS.
GO; GO:0006955; P:immune response; NAS.
InterPro; IPR003599; Ig.
InterPro; IPR003596; Ig.
InterPro; IPR013106; V-set.
InterPro; IPR013106; V-set.
Pfam; PF07686; V-set; 1.
SMART; SM00409; IG; 1.
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P06314;
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Distributed under the Creative Commons Attribution-NoDerivs License
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Mammalia, Eutheria, Euarchontoglires, Primates, Catarrhini, Hominidae
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                                                                     Ig kappa chain V-IV region B17.
/FTId=PRO_000015183.
Framework-1.
Complementarity-determining-1.
Framework-2.
Complementarity-determining-2.
Framework-3.
Complementarity-determining-2.
Framework-4.
By similarity.
                                                                                                                                                                                                                                                                        Length 134;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    integrated into UniProtKB/Swiss-Prot
                                                                                                                                                                                                                                                                     Score 454; DB 1; I Pred. No. 2.1e-40; 3; Mismatches 15;
                                          region;
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PIR; A01904; K4HUJI.
HSSP; P01625; 1LVE.
SMR; P06313; 21-133.
GO; GO:0005576; C:extracellular region; NAS.
GO; GO:0005823; F:antigen binding; NAS.
GO; GO:0006955; P:immune response; NAS.
InterPro; IPR003599; Ig.
InterPro; IPR007110; Ig-like.
InterPro; IPR013106; V-set.
InterPro; IPR013106; V-set.
SMART; SM00409; IG; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
Immunoglobulin domain; Immunoglobulin V region SIGNAL
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                                        Immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUCLEOTIDE SEQUENCE [GENOMIC DNA]
MEDLINE=86041853; PubMed=2997712;
Klobeck H.G., Bornkamm G.W., Comb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-JAN-1988, sequence version 1
07-MAR-2006, entry version 43.
Ig kappa chain V-IV region JI p.
Homo sapiens (Human).
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14966 MW;
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        IGv; 1.
5; IG_LIKE;
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SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIK
Immunoglobulin domain; I
SIGNAL
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86; Conserv
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NCBI_TaxID=9606;
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Distributed under the Creative Commons Attribution-NoDerivs License
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Primates; Catarrhini; Hominidae
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/FTId=PRO_000015182.
Framework-1.
Complementarity-determining-1.
Framework-2.
Complementarity-determining-2.
Framework-3.
Complementarity-determining-3.
Framework-4.
By similarity.
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HGNC; HGNC:5834; IGKV4-1.

GO; GO:0005576; C:extracellular region; NAS.

GO; GO:0003823; F:antigen binding; NAS.

GO; GO:0006955; P:immune response; NAS.

InterPro; IPR003599; Ig.

InterPro; IPR003599; Ig.

InterPro; IPR003596; Ig.

InterPro; IPR013106; V-set.

Pfam; PF07686; V-set; 1.

SMART; SM00409; IG; 1.

SMART; SM004009; IG; 1.

SMART; SM00406; IGV; 1.

PROSITE; PS50835; IG LIKE; 1.

Immunoglobulin domain; Immunoglobulin V region; Signature of the state of the s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         one Ig kappa V-IV
                                                                                                                                                                                                                                                                                                                                                                                                        988, integrated into UniProtKB/Swiss-Prot
988, sequence version 1.
006, entry version 45.
chain V-IV region precursor (Fragment).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUCLEOTIDE SEQUENCE [GENOMIC DNA].
MEDLINE=86041853; PubMed=2997712;
Klobeck H.G., Bornkamm G.W., Combriato (Zachau H.G.;
"Subgroup IV of human immunoglobulin K single germline gene.";
Nucleic Acids Res. 13:6515-6529(1985).
-!- MISCELLANEOUS: There is only one Ig
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No
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Mammalia; Eutheria; Euarchontoglires;
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PIR; A01902; K4HU.
HSSP; P01625; 1LVE.
SMR; P06312; 21-121.
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Name=IGKV4-1;
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01-JAN-1988,
01-JAN-1988,
07-MAR-2006,
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Distributed under the Creative Commons Attribution-NoDerivs License
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Glires; Rodentia; Sciurognathi;
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Framework-1.
Complementarity-determining-1.
Framework-2.
Complementarity-determining-2.
Framework-3.
Complementarity-determining-3.
By similarity.
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Framework-1.
Complementarity-determining-1
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                                                                                                                                                                        Length
                                                                                                                                          CRC64
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              21-JUL-1986, integrated into UniProtKB/Swiss-Prot.
21-JUL-1986, sequence version 1.
07-MAR-2006, entry version 50.
Ig kappa chain V-V region MOPC 21 precursor.
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodent Muroidea; Muridae; Murinae; Mus.
NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                            Score 410; DB 1; Pred. No. 9.6e-36; D; Mismatches 14;
                                                                                                                                        9586AD4188D33974
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PIR; A93736; KVMS21.
PDB; 11GC; X-ray; L=30-136.
Ensembl; ENSMUSG0000060300; Mus musculus.
LinkHub; P01634; -.
InterPro; IPR003599; Ig.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig.
InterPro; IPR013106; V-set.
Pfam; PF07686; V-set; 1.
SMART; SM00409; IG; 1.
SMART; SM00409; IG; 1.
SMART; SM00406; IGv; 1.
SMART; SM00406; IGv; 1.
SMART; SM00406; IG LIKE; 1.
SMART; SM00406; IG LIKE; 1.
SMART; SM010406; IG LIKE; 1.
SMART; SM00406; IG LIKE; 1.
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MEDLINE=82059477; PubMed=6170937;
Hamlyn P.H., Gait M.J., Milstein C.;
"Complete sequence of an immunoglobulin and the dideoxynucleotide method of RNA Nucleic Acids Res. 9:4485-4494(1981).
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MEDLINE=73053310; PubMed=4638343;
Svasti J., Milstein C.;
"The complete amino acid sequence
Biochem. J. 128:427-444(1972).
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76.58;
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P01632;
21-JUL-1986, i
21-JUL-1986, E
07-MAR-2006, e
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TISSUE=Abdominal adipose tissue;

MEDLINE=98249779; PubMed=9588180; DOI=10.1006/bbrc.1998.8515;

Olsen K.E., Sletten K., Westermark P.;

"Extended analysis of AL-amyloid protein from abdominal wall subcutaneous fat biopsy: kappa IV immunoglobulin light chain.";

Biochem. Biophys. Res. Commun. 245:713-716(1998).

-!- FUNCTION: May play an important role in fibrillogenesis.

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                                                                                                                                                                                                                                                                                                            30-MAY-2003, integrated into UniProtKB/Swiss-Prot.
30-MAY-2003, sequence version 1.
07-MAR-2006, entry version 23.
Ig kappa chain V-IV region STH (Fragment).
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                       -- TYVSWYQQKPEQSPKLLIYGASN
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   Framework-2.
Complementarity-determining-2
Framework-3.
Complementarity-determining-3
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No. 8e-35;
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Pred. No. 8e-3.
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GNIVMTQSPKSMSMSVGERVTLTCKASENVV
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InterPro; IPR003599;
InterPro; IPR007110;
InterPro; IPR003596;
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    P83593;
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HSSP; Q9ER29; ZAP2.
SMR; P01632; 1-114.
Ensembl; ENSMUSG0000052500; Mus musculus.
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MEDLINE=81241357; PubMed=6788890; DOI=10.1084/jem.153.5.1366;
Kwan S.-P., Rudikoff S., Seidman J.G., Leder P., Scharff M.D.;
"Nucleic acid and protein sequences of phosphocholine-binding light
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antibody response to
lot mutated T15 antibodies
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurogna;
Muroidea; Muridae; Murinae; Mus.
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Framework-2.
Complementarity-determining-2
Framework-3.
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Complementarity-determining-
Framework-4.
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No. 7.8e-35;
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Mismatches
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J. Immunol. 141:4012-4019(1988).
-!- FUNCTION: Anti-phosphocholine antibody
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Immunoglobulin
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MEDLINE=89035545; PubMed=3141511;
Claflin J.L., Berry J.;
"Genetics of the phosphocholine-specific Streptococcus pneumoniae. Germ-line but r
                                                                                                                                                                                                                                                                                                                                                                                                Score
Pred.
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12060 MW;
InterPro; IPR013106; V-set.
Pfam; PF07686; V-set; 1.
SMART; SM00409; IG; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1
Direct protein sequencing; Immunoglobulin V region.
CHAIN 1 >109
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70.6%;
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NCBI_TaxID=10090;
[1]
NUCLEOTIDE SEQUENCE.
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MEDLINE=22716456; PubMed=12833571; DOI=10.1002/jmr.617;
Erlandsson A., Holm P., Ullen A., Stigbrand T., Sundstrom B.E.;
"Studies of the interactions between the anticytokeratin 8 monocleantibody TS1, its antigen and its anti-idiotypic antibody alphaTS:
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Framework-2.
Complementarity-determining-2.
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Framework-4.
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MGI; MGI:3577282; U29423.

InterPro; IPR003599; Ig.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
InterPro; IPR013106; V-Set.

R SMART; SM00409; IG; 1.
R SMART; SM00406; IGv; 1.
R PROSITE; PS50835; IG_LIKE; 1.
R PROSITE; PS50835; IG_LIKE; 1.
Immunoglobulin domain; Immunoglobulin V-I regic /FrId=PRO_0000059770.

"HAIN I >114 Ig kappa chain V-I regic /FrId=PRO_0000059770.
Framework-1.
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Pred. No. 1.7e
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InterPro; IPR003599; Ig.
InterPro; IPR007110; Ig-like.
InterPro; IPR013596; Ig_v.
InterPro; IPR013106; V-Bet.
SMART; SM00409; IG; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
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15-MAR-2005, integrated into Unip 15-MAR-2005, sequence version 1.

07-FEB-2006, entry version 6.

Kappa light chain variable region Name=IgG1 anti-TS1 VL;

Mus musculus (Mouse).
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12717 MW;
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68.1%;
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21-JUL-1986, sequence version 1.
07-MAR-2006, entry version 45.
Ig kappa chain V-V region MPC11 precursor.
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi, Muroidea; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "Sequence of the full-length immunoglobulin kappa-chain of mouse myeloma MPC 11.";
Biochem. J. 171:337-347(1978).

-!- MISCELLANEOUS: The mature chain has 12 additional residues at amino end, due to a tandem duplication of 36 nucleotides after codon for residue 36. Residue 42 corresponds to the N-terminal residue of typical kappa chains.
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MEDLINE=80176554; PubMed=6245773;
Rabbitts T.H., Hamlyn P.H., Matthyssens G., Roe B.A.;
"The variability, arrangement, and rearrangement of immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUCLEOTIDE SEQUENCE [GENOMIC DNA] OF 1-71.
MEDLINE=83001944; PubMed=6288267; DOI=10.1016/0092-8674(82)90184
Kelley D.E., Coleclough C., Perry R.P.;
"Functional significance and evolutionary development of the 5'-terminal regions of immunoglobulin variable-region genes.";
Cell 29:681-689(1982).
                                                                                                                                                                                                                                                                                N
                                                                                                        Length 115
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                                                              CRC64
                                                                              391; DB 2; L6
No. 9.9e-34;
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Can. J. Biochem. 58:176-187(1980)
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MEDLINE=78186617; PubMed=418775
Smith G.P.;
                                                                                                                                                    14;
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HSSP; P01634; 11GC.
SMR; P01633; 42-149.
Ensembl; ENSMUSG0000061362; MINTERPRO; IPR003599; IG.
InterPro; IPR003599; IG.
InterPro; IPR003596; Ig.
InterPro; IPR003596; Ig.
InterPro; IPR013106; V-Set.
Pfam; PF07686; V-Set; 1.
SMART; SM00409; IG; 1.
SMART; SM00406; IGV; 1.
                                                             12560 MW;
                                                                                                        64.5%;
ilarity 68.4%;
Conservative 1
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domain
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
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MEDLINE=93170387; PubMed=8436174;
Wagner S.D., Luzzatto L.;
"V kappa gene segments rearranged in chronic lymphocytic leukemia distributed over a large portion of the V kappa locus and do not segmentic mutation.";
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MEDLINE=98277139; PubMed=9614934; DOI=10.1006/clin.1998.4531;
Wu X., Liu B., Van der Merwe P.L., Kalis N.N., Berney S.M.,
                                            g kappa chain V-V region MPC11
FTId=PRO_000015189.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUCLEOTIDE SEQUENCE.
MEDLINE=92202880; PubMed=1552291; DOI=10.1084/jem.175.4.983;
Martin T., Duffy S.F., Carson D.A., Kipps T.J.;
"Evidence for somatic selection of natural autoantibodies.";
J. Exp. Med. 175:983-991(1992).
                                                                                                                                                                                                                                                                   149;
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                                                                                                                                                      Complementarity-determining-2 Framework-3.
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                                                                                                           Framework-1.
Complementarity-determining
Framework-2.
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Complementarity-determining
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D.R., Lerner
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   domain;
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                                                                                                                                                                                                                                                                 DB 1,
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Pred. No. 6e-3.
2; Mismatches
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  Immunoglobulin peat; Signal.
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MEDLINE=92228746; PubMed=1373487;
Zebedee S.L., Barbas C.F. 3rd, Hom Y.L.,
DeGraw J., Pyati J., LaPolla R., Burton I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-MAY-2000, sequence version 1.
07-FEB-2006, entry version 22.
Myosin-reactive immunoglobulin light
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Wu X., Liu B., Van der Merwe P.L.,
Young D.C.;
"Myosin-reactive autoantibodies in
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23:391-397(1993)
                Repeat;
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16434 MW;
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ilarity 66.7%;
Conservative 12
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 Wirect protein sequencing;
mmunoglobulin V region; RiGNAL 1 29
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76; Conserv
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                                                                                                                                                                                                                               SEQUENCE
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Catarrhini; Hominidae;
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NUCLEOTIDE SEQUENCE.
MEDLINE=88171307; PubMed=3127527; DOI=10.1084/jem.167.3.840;
Kipps T.J., Tomhave E., Chen P.P., Carson D.A.;
Kipps T.J., Tomhave E., Chen P.P., Carson D.A.;
"Autoantibody-associated kappa light chain variable region ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
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Chordata; Craniata;
Euarchontoglires; P
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P18135;
01-NOV-1990, integrated into
01-NOV-1990, sequence version
07-MAR-2006, entry version 4.
Ig kappa chain V-III region 1.
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata
Mammalia; Eutheria; Euarchon
                                AAD56272.1;
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64.6%;
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LIKE;
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                         EMBL; AF035036; AAD56272
PIR; A30601; A30601.
PIR; A30601; B30601.
PIR; B30601; B30601.
PIR; B30601; B30601.
PIR; C30601; C30601.
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PIR; C30601; D30601.
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PIR; D30601; D30601.
PIR; B30601; B30608.
PIR; B30601; B30608.
PIR; B30601; B30608.
PIR; B30608; B30608.
PIR; B44151; H44151.
PIR; PH0963; PH0963.
PIR; PH0963; PH0963.
PIR; PH0964; PH0964.
PIR; PH0965; PH0965.
PIR; PH0965; PH0966.
InterPro; IPR003599; IglinterPro; IPR003596; IglinterPro; IPR003596; IglinterPro; IPR003596; IglinterPro; IPR003596; V-8
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InterPro; IPR0035,
InterPro; IPR013106; V-SMART; SM00409; IG; 1.
SMART; SM00406; IGv; 1
PROSITE; PS50835; IG I
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                                                                                               see http://www.uniprot.org/terms
Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Muridae; Murinae; Mus.
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     somatic
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STRAIN=CZECH II; TISSUE=Mammary tumor metastatized to lung.
MMTV-LTR/Wnt1 model. Expression driven by an MMTV-LTR enhancer.;
MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
                                                         lymphocyti
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                                             immunoglobulin
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JK1 segment.
By similarity.
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Complementarity-determining-Framework-3.
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expressed in chronic lymphocytic leukemia with little mutation. Implications for etiology and immunotherapy. J. Exp. Med. 167:840-852(1988).

-1- DISEASE: The protein is one of the surface immunog autoantibodies expressed in patients with chronic leukemia.
                                                                                                                                                                                                                                                                                                                                                                                                   chain V-III
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Pred. No. 1.2e-32;
7; Mismatches 19;
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PIR; PL0022; K3HUHA.

HSSP; P01625; LEEQ.

SMR; P18135; 21-129.

LinkHub; P18135; 21-129.

R GO; GO:0003823; F:antigen binding; NAS.

R GO; GO:0006955; P:immune response; NAS.

DR InterPro; IPR003599; Ig.

InterPro; IPR00110; Ig-like.

DR InterPro; IPR001306; V-set.

DR SMART; SM00409; IG; 1.

DR SMART; SM00406; IGv; 1.

DR SMART; SM00406; IGv; 1.

PROSITE; PS50835; IG_LIKE; 1.

Tmmunoglobulin domain; Immunoglobulin V

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Framework-

Pramework-

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64.0%;
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A Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

A Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

A Richards S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

A Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

A Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

A Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,

Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,

A Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;

A "Generation and initial analysis of more than 15,000 full-length human
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Distributed under the Creative Commons Attribution-NoDerivs License
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P04207;
20-MAR-1987, integrated into UniProtKB/Swiss-Prot.
01-NOV-1990, sequence version 2.
07-MAR-2006, entry version 43.
Ig kappa chain V-III region CLL precursor (Rheumatoid factor).
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae.
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STRAIN=CZECH II; TISSUE=Mammary tumor metastatized to lung.
MMTV-LTR/Wnt1 model. Expression driven by an MMTV-LTR enhancer.;
NIH MGC Project;
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Submitted (MAR-2005) to the EMBL/GenBank/DDBJ databases
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Pred. No. 2.5e-32;
1; Mismatches 24;
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MGI; MGI:96495; Igk-C.
InterPro; IPR003599; Ig.
InterPro; IPR0031006; Ig_MHC.
InterPro; IPR003596; Ig_V.
InterPro; IPR003596; Ig_V.
InterPro; IPR013106; V-Set.
Pfam; PF07654; C1-Set; 1.
SMART; SM00409; IG; 1.
SMART; SM00406; IGV; 1.
PROSITE; PS50835; IG_LIKE; 2.
PROSITE; PS00290; IG_MHC; UNKNOWN 1.
SEQUENCE 239 AA; 26302 MW; 98FC4
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NUCLEOTIDE SEQUENCE [GENOMIC DNA].

MEDLINE=86177570; PubMed=3083417;

Jirik F.R., Sorge J., Fong S., Heitzmann J.G., Curd J.G., Chen P.P.,
Goldfien R., Carson D.A.;
"Cloning and sequence determination of a human rheumatoid factor
light-chain gene.";
Proc. Natl. Acad. Sci. U.S.A. 83:2195-2199(1986).

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Framework-1.
Complementarity-determining-1.
Framework-2.
Complementarity-determining-2.
Framework-3.
Complementarity-determining-3.
JKl segment.
By similarity.
                                                                                                                                                                                                                                                   EMBL; M12740; AAA58992.1; -; Genomic_DNA.
HSSP; P01625; 1LVE.
SMR; P04207; 21-129.
LinkHub; P04207; -.
GO; GO:0003823; F:antigen binding; NAS.
GO; GO:0006955; P:immune response; NAS.
InterPro; IPR003599; Ig.
InterPro; IPR003106; V-set.
InterPro; IPR003596; Ig-v.
InterPro; IPR013106; V-set.
Pfam; PF07686; V-set; 1.
SMART; SM00409; IG; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
Immunoglobulin domain; Immunoglobulin V region; Signal.
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Pred. No. 1.5e-32;
21; Mismatches 14;
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14275 MW;
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Best Local Similarity 63.5%;
Matches 73; Conservative 2
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-936A-8

. CQNDHIYPYTFGGGTKLEIK SRGDIVMTQSPSSLSVSAGE US-09-889-606 core Sequence Title: Perfect

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residues 87530628 seds, 650591 Searched

6505 parameters: chosen satisfying hits of number Total

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Match 0% Match 100% First 45 su Minimum Maximum Listing Post-processing:

summaries

Databas

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/EMC\_Celerra\_SIDS3/ptodata/2/iaa/5\_COMB.pep:\*
/EMC\_Celerra\_SIDS3/ptodata/2/iaa/6\_COMB.pep:\*
/EMC\_Celerra\_SIDS3/ptodata/2/iaa/H\_COMB.pep:\*
/EMC\_Celerra\_SIDS3/ptodata/2/iaa/H\_COMB.pep:\*
/EMC\_Celerra\_SIDS3/ptodata/2/iaa/PCTUS\_COMB.pep:\*
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		ď			SUMMARIES	
Result		Query				
No.	SC	Match	Length	DB	ΩI	Description
<b>-</b>	54		113	-	US-08-467-420A-45	ence 45, App
7	4	89.3	113	Н	US-08-470-110A-45	ence 45, App
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4	4	89.3	113	-	US-08-940-371-45	ence 45, App
ស	4	89.3	113	7	US-08-637-647-45	equence 45, App
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80	$\sim$	88.4	113	-	US-08-077-252B-4	4
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12	536	88.4	113	-	US-08-082-842A-2	Sequence 2, Appli
13	m	88.4	러	~	US-09-002-753A-4	4,
14	$\mathbf{c}$	88.4	Н	N	-09-657-27	quence 4,
15	C	88.4	113	ഗ	9	quence 4,
16	3	88.4	115	<b>⊢</b>	US-07-942-245-7	ence 7,
17	ന	88.3	113	Н	-08-467-420A-4	quence 46
18	m	88.3	113	٦	US-08-470-110A-46	ence 46
19	3	· :	113	Н	US-08-667-769A-46	ence 46
20	m	· : ω	113	႕	US-08-940-371-46	ence 46
21	m	88.3	113	7	US-08-637-647-46	46
22	ന	88.3	113	~	US-10-700-740-46	ence 46
23	m	88.3	113	ហ	PCT-US95-17082A-46	ence 46
24	ന	88.1	277	Н	US-08-256-790-2	equence 2,
25	N	87.1	113	٦	디	equence 12
56	O	87.1	113	~	US-08-765-469-12	12

equence 26, A equence 16, A equence 16, A	equence 16, equence 16, equence 47, equence 47,	equenc equenc equenc equenc		equence 4, A equence 28, equence 21,
09-232-290-26 08-667-769A-16 10-700-740-16	T-US95-17082 -09-509-031- -08-442-542-	09-561-500 09-561-108 09-561-526 09-561-499	US-09-998-831-9 US-09-561-005-9 US-09-562-245-9 US-08-513-968-36	-09-587-526-4 -09-393-627B-
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## ALIGNMENTS

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GENERAL INFORMATION:
APPLICANT: Ames, Sobert S.
APPLICANT: Ames, Stobert S.
APPLICANT: Chaiken, Irwin M.
APPLICANT: Chaiken, Irwin M.
APPLICANT: Coss, Mitchell S.
APPLICANT: Goos, Mitchell S.
APPLICANT: Goos, Mitchell S.
APPLICANT: Holmes, Stephen D.
APPLICANT: Holmes, Stephen D.
APPLICANT: Theisen, Timothy W.
TITLE OF INVENTION: Treatment of IL5 Mediated Disorders
NUMBER OF SEQUENCES: 74
CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corp./Corporate
ADDRESSEE: Intellectual Property
STREET: P. O. Box 1539-UW220
CITY: King of Prussia
STATE: Pennsylvania
COWNPUTER: Dennsylvania
COWNPUTER: EADABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURSTREAT PAPLICATION DATA:
APPLICATION NUMBER: US/08/467,420A
FILING DATE:
CLASSIFICATION NUMBER: US 08/363131
FILING DATE: 23-DEC-1994
ATTORNEY APPRORENTION:
NAME: SHIFTM: AFFERNATION:
NAME: SHIFTM: THE TEAD ATTORNEY AND ADDRESSEE
NAME: SHIFTM: THE TEAD ADDRESSEE
CLASSIFICATION NUMBER: US 08/363131
FILING DATE: 23-DEC-1994
ATTORNEY AGENT INFORMATION:
NAME: SHIFTM: THE TEAD ADDRESSEE
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REGISTRATION NUMBER: 34,028

REFERENCE/DOCKET NUMBER: P50282

TELEPHONE: 610 270-5024

TELEFAX: 610 270-5090

INFORMATION FOR SEQ ID NO: 45:

SEQUENCE CHARACTERISTICS:

LENGTH: 113 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear
RESULT 1
US-08-467-420A-45
; Sequence 45, Application US/08467420A
; Patent No. 5683892
; GENERAL INFORMATION:
; APPLICANT: Ames, Robert S.
; APPLICANT: Appelbaum, Edward R.
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US/08/667,769
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ER: P50503
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APPLICANT: Appelbaum, Edward R.
APPLICANT: Chaiken, Irwin M.
APPLICANT: Cook, Richard M.
APPLICANT: Gross, Mitchell S.
APPLICANT: Holmes, Stephen D.
APPLICANT: McMillan, Lynette J.
APPLICANT: Theisen, Timothy W.
TITLE OF INVENTION: Treatment of
                                                                                                                                                ce 45, Application US/08667769A
No. 5783184
AL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                              Beecham
-UW2220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-270-5024
TELEFAX: 610-270-5090
INFORMATION FOR SEQ ID NO: 45:
SEQUENCE CHARACTERISTICS:
LENGTH: 113 amino acids
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ilarity 91.2%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: SmithKline
STREET: P.O. Box 1539-
CITY: King of Prussia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: USE FILING DATE: CLASSIFICATION: 424
PRIOR APPLICATION DATA: APPLICATION NUMBER: PO
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CORRESPONDENCE ADDRESS:
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                                                                                                                    RESULT 3
US-08-667-769A-45
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ILS Mediated Disorders
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                                            , DB 1;
2.2e-42;
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Pred. No. 2.2e-42
                                           Score 541; DB Pred. No. 2.2e 4; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 74
CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corp
ADDRESSEE: Intellectual Property
STREET: P. O. Box 1539-UW2220
CITY: King of Prussia
STATE: Pennsylvania
COUNTRY: USA
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COUNTRY: USA
ZIP: 19406-0939
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, V
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34,028
50282
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Patent No. 5693323
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                              ICANT: Ames, Robert S.
ICANT: Appelbaum, Edward R.
ICANT: Chaiken, Irwin M.
ICANT: Cook, Richard M.
ICANT: Gross, Mitchell S.
ICANT: Holmes, Stephen D.
ICANT: Holmes, Stephen D.
ICANT: Theisen, Timothy W.
ICANT: Theisen, Timothy W.
E OF INVENTION: Recombinant I
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APPLICATION NUMBER: US 08/3631
FILING DATE: 23-DEC-1994
ATTORNEY/AGENT INFORMATION:
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TELEPHONE: 610 270-5024
TELEFAX: 610 270-5090
FORMATION FOR SEQ ID NO: 45:
                                         89.3%;
ilarity 91.2%;
Conservative
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91.2%;
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REGISTRATION NUMBER: 34
REFERENCE/DOCKET NUMBER:
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TOPOLOGY: linear
LECULE TYPE: pro
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les 103; Conser
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nes 103; Conser
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CLASSIFICATION:
TYPE:
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US-08-467-420A-45
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APPLICANT:
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Mediated Disorders
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2.2e-42;
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of ILS
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/470110
FILING DATE: 06-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/467420
FILING DATE: 06-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/363131
FILING DATE: 23-DEC-1994
                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 19406-0939

ZIP: 19406-0939

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, V
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Sequence 45, Application US/10700740
Patent No. 6946130
GENERAL INFORMATION:
APPLICANT: Ames, Robert S.
                              ESULT 5
S-08-637-647
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Best Loc
Matches
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DIVMTQSPSSLSVSAGEKVTMSCKSSQSLLNSRNQKNYLAWYQQKPGQPPKLLIYGVFIR
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Intellectual Property
O. Box 1539-UW2220
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ed. No. 2.2e-42;
Mismatches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA

ZIP: 19406-0939

ZIP: 19406-0939

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, V

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/940,371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US/08/470,110
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Pred.
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                                                                                                                                                                                      APPLICANT: Ames, Robert S.
APPLICANT: Appelbaum, Edward R.
APPLICANT: Chaiken, Irwin M.
APPLICANT: Cook, Richard M.
APPLICANT: Gross, Mitchell S.
APPLICANT: Holmes, Stephen D.
APPLICANT: Holmes, Stephen D.
APPLICANT: Theisen, Timothy W.
TITLE OF INVENTION: Recombinant I
TITLE OF INVENTION: Treatment of
NUMBER OF SEQUENCES: 74
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                P50282
                                                                                                                                               ce 45, Application US/08940371
No. 5851525
AL INFORMATION:
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TELEPHONE: 610 270-5024
TELEFAX: 610 270-5090
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ilarity 91.2%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
APPLICATION NUMBER: US 0
FILING DATE: 23-DEC-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Sutton, Jeffrey A. REGISTRATION NUMBER: 34, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: SmithKline HADDRESSEE: Intellectual STREET: P. O. Box 1539-CITY: King of Prussia STATE: Pennsylvania
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 113 amino acid
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CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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amino acid
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Best Local Similarity
Matches 103; Conser
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TOPOLOGY: lin
MOLECULE TYPE:
8-940-371-45
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US-08-940-371-45
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                                                                                                                                             Sequence 45,
Patent No. 58
GENERAL INFC
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                                                                                                                                             ILS Antagonists Useful
ILS Mediated Disorders
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                                                                                                                                                                                                        Corp./Corporate
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Pred. No. 2.2e-42
; Mismatches
                                                                                                                                                                                                                                                                                                                                                                  Version
                                                                                                                                       TITLE OF INVENTION: Recombinant IL5 AU
TITLE OF INVENTION: Treatment of IL5 N
NUMBER OF SEQUENCES: 50
CORRESPONDENCE ADDRESS:
ADDRESSEE: Smithkline Beecham Corp.,
ADDRESSEE: Intellectual Property
STREET: P. O. Box 1539-UW2220
CITY: King of Prussia
STATE: Pennsylvania
COUNTRY: USA
ZIP: 19406-0939
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: PatentIn Release #1.0, Ven
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/637,647
FILING DATE: 19-JUN-1997
CLASSIFICATION NUMBER: US/08/363,131
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/363,131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Sutton, Jeffrey A.
REGISTRATION NUMBER: 34,028
REFERENCE/DOCKET NUMBER: P50282
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610 270-5090
                                        Ames, Robert S.
Appelbaum, Edward R.
Chaiken, Irwin M.
Cook, Richard M.
Gross, Mitchell S.
McMillan, Lynette J.
Theisen, Timothy W.
NVENTION: Recombinant
NVENTION: Treatment of
Sequence 45, Application US/0863764 Patent No. 6129913 GENERAL INFORMATION:
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ilarity 91.2%;
Conservative
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CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 113 amino acid
TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protein
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STRANDEDNESS: 811
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nes 103; Conser
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INFORMATION FOR
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                                                                                   APPLICANT:
APPLICANT:
APPLICANT:
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APPLICANT:
                                                                        APPLICANT:
                                                                                                                                 APPLICANT:
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                                                                                                                                                               COUNTRY: USA
ZIP: 19406-0939
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MOLECULE TYPE:
IS95-17082A-45
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-077-252B-4
    APPLICANT:
APPLICANT:
APPLICANT:
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                                                                                                Antagonists Useful
Mediated Disorders
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                                                                                                                                                                                                              ZIP: 19406-0939

ZIP: 19406-0939

ZIP: 19406-0939

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC Compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/700,740

FILING DATE: 04-No. 6946130-2003

CLASSIFICATION DATA:

APPLICATION NUMBER: US/10/144,644

FILING DATE: 06-JUN-1995

APPLICATION NUMBER: US 08/467420

FILING DATE: 06-JUN-1995

APPLICATION NUMBER: US 08/467420

FILING DATE: 23-DEC-1994
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Pred. No. 2.2e-42;
!; Mismatches 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Sutton, Jeffrey A.
REGISTRATION NUMBER: 34,028
REFERENCE/DOCKET NUMBER: P50282-2
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-270-5024
TELEFAX: 610-270-5090
INFORMATION FOR SEQ ID NO: 45:
SEQUENCE CHARACTERISTICS:
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of ILS
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ADDRESSEE: SmithKline Beecham
STREET: P.O. Box 1539-UW2220
CITY: King of Prussia
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PCT-US95-17082A-45
Sequence 45, Application PC/TUS9517082A
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL TOTAL AMES, Robert S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ID NO:
                                                                                                INVENTION: Recombinant Treatment o
Appelbaum, Edward R
Chaiken, Irwin M.
Cook, Richard M.
Gross, Mitchell S.
Holmes, Stephen D.
McMillan, Lynette J
Theisen, Timothy W.
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TOPOLOGY: linear
CULE TYPE: protein
ENCE DESCRIPTION: SEQ
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Appelbaum, Edward I
Chaiken, Irwin M.
Cook, Richard M.
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                                                                                                                                                                             CITY: King of Pruss
STATE: Pennsylvania
COUNTRY: USA
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ilarity 91.2%;
Conservative
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amino acid
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-10-700-740-45
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APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                           NUMBER
                                                                                                  TITLE
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Fragments Having Binding Specificity
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                                                                    of ILS Antagonists Useful of ILS Mediated Disorders
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                                                                                                                                        Corp./Corporat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 541; DB 5;
Pred, No. 2.2e-42;
Mismatches
                                                                                                                                                                                                                                                                                                                 Version
                                                                                                                                                                                                                                                                                                                                                 PCT/US95/17082A
                                                                                                                                                                                                                                        COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, V
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/17082
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MBER: US 08/467420
06-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US 08/363131
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ER: P50282-
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APPLICANT: Pastan, Ira
APPLICANT: Lee, Byungkook
APPLICANT: Jung, Sun-Hee
APPLICANT: Brinkmann, Ulrich
TITLE OF INVENTION: Recombinant Di
TITLE OF INVENTION: Polypeptide Fi
NUMBER OF SEQUENCES: 23
                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US 08/47011
FILING DATE: 06-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/46742
                                                                   Recombinant
Treatment of
76
tchell S.
tephen D.
Lynette J.
                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham
STREET: P.O. Box 1539-UW2220
CITY: King of Prussia
STATE: Pennsylvania
Gross, Mitchell S.
Holmes, Stephen D.
McMillan, Lynette J.
Theisen, Timothy W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 23-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: Sutton, Jeffrey A.
REGISTRATION NUMBER: 34,028
REFERENCE/DOCKET NUMBER: P50
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-270-5024
TELEFAX: 610-270-5024
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Best Local Similarity 91.2%;
Matches 103; Conservative
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                                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 06-JUN-
NIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein
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                                                  APPLICANT: Theisen, T
TITLE OF INVENTION: F
TITLE OF INVENTION: T
NUMBER OF SEQUENCES:
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/note= "Light chain variable region Pro residue that can be changed to Cys, paired with Heavy chain variable region position 114 Ala residue change to Cys, for possible S2 site interchain disulfide bond"
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/note= "Complementarity Determining
Region 1 (CDR1)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    - CDR2
"Complementarity Determining
2 (CDR2)"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           oclonal antibody variable region
                                                   l and Crew LLP
Eighth Floor
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"Framework Region
                                                                                                                                           COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/077,252B
FILING DATE: 14-JUN-1993
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen Lauver
REGISTRATION NUMBER: 15280-152
REFERENCE/DOCKET NUMBER: 15280-152
REFERENCE/DOCKET NUMBER: 15280-152
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 113 amino acid
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /label= FR1
/note= "Framework Region
                                                                                                                                                                                                                       #1
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Light chain variab
                                                   Townsend
Center, E
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/note= "
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/note= "
Region 2
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SSEE: Townsend and T
T: Two Embarcadero Co
San Francisco
: California
Y: USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: protein FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: Region
LOCATION: 41..55
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: Modified LOCATION: 49
OTHER INFORMATION: OTHER INFORMATION: OTHER INFORMATION: OTHER INFORMATION: OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: Protein
LOCATION: 1..113
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: Region
LOCATION: 24.40
OTHER INFORMATION:
OTHER INFORMATION:
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LOCATION: 56..62
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: Region
LOCATION: 63..94
OTHER INFORMATION:
OTHER INFORMATION:
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LOCATION: 1.23
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
                                CORRESPONDENCE ADDRESSEE: To STREET: Two I CITY: San Fre STATE: Califo COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 113
TYPE: amino
STRANDEDNESS:
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V-H97
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             /note= "V-L93 Ser to Tyr mutation (pseudo-symmetrically related to Ser to Tyr mutation site in Heavy chain variable region)"
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Pred. No. 6.3e-42
); Mismatches
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N.W., Suite
                                                                                                                                                                                                        /label= FR4
/note= "Framework
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: PCT/US92/10906
FILING DATE: 14-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/808,464
FILING DATE: 13-DEC-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne, Kessler, Golds'
STREET: 1100 New York Ave., N.W.,
CITY: Washington
STATE: D.C.
COUNTRY: United States of America
ZIP: 20005-3934
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         America
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US-08-107-669D-2
; Sequence 2, Application US/08107669D
; Patent No. 5766886
; GENERAL INFORMATION:
; APPLICANT: Studnicka, Gary M.
; TITLE OF INVENTION: Modified Antib
; TITLE OF SEOUENCES: 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/107,66
FILING DATE: 13-AUG-1993
CLASSIFICATION: 424
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REGISTRATION NUMBER: 33,851
                                                                                                                                              Region
                                                                                                                  /label=
/note=
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ilarity 91.2%;
Conservative
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APPLICATION DATA:
APPLICATION NUMBER: P(
FILING DATE: 14-DEC-19
                                                                                                                                                                            NAME/KEY: Region
LOCATION: 104..109
OTHER INFORMATION:
OTHER INFORMATION:
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                                                                                     NAME/KEY: Region
LOCATION: 95..103
OTHER INFORMATION:
OTHER INFORMATION:
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OTHER INFORMATION: OTHER INFORMATION: OTHER INFORMATION: OTHER INFORMATION: OTHER INFORMATION: OTHER INFORMATION:
OCATION: 93
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OPERATING SYSTEM:
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Best Local Similarity
Matches 103; Conser
                                                                                                                                                                          NAME/KEY:
LOCATION:
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US-08-477-531B-2
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Matches 103
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STATE:
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N.W., Suite 600
                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Studnicka, Gary M.
TITLE OF INVENTION: Modified Antibody Variable Domains
NUMBER OF SEQUENCES: 89
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne, Kessler, Goldstein and Fox P.L.L.(
STREET: 1100 New York Ave., N.W., Suite 600
CITY: Washington
                                                                                                                                                     Score 536; DB 1; Le
Pred. No. 6.3e-42;
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   0610.1000001/MAC
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R: 0610.1000003
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APPLICATION NUMBER: 08/082,842
FILING DATE: 23-JUN-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/10906
FILING DATE: 14-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/808,464
FILING DATE: 13-DEC-1991
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                             Sequence 2, Application US/08472788A
Patent No. 5770196
GENERAL INFORMATION:
APPLICANT: Studnicka, Gary M.
REFERENCE/DOCKET NUMBER: 0610
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202/371-2600
TELEFAX: 202/371-2600
TELEFAX: 202/371-2540
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 113 amino acids
TYPE: amino acids
TYPE: amino acid
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91.2%;
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NAME: Cimbala, Michele A
REGISTRATION NUMBER: 33,
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202/371-2540
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LENGTH: 113 amino acide
                                                                                                                                                                                             Conservative
                                                                                                                          protein
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ZIP: 20005-3934
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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Best Local Similarity
Matches 103; Conser
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TYPE:
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 . DB 1;
6.3e-42;
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Pred. No. 6.3e-42;
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 33,851
REFERENCE/DOCKET NUMBER: 0610.1000004/MAC
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202/371-2540
TELEFAX: 202/371-2540
                                                                                                                                                                                                                                                                                                                                                                                   Version
ore 536; DB
ed. No. 6.3e
Mismatches
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APPLICATION NUMBER: US 0.7 / ATTORNEY/...
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N: 436
                                                                                                                                                                                                                                                                                                               America
                                                                                                                                                                                                                                                                                                                                                           IBM PC compatible
YSTEM: PC-DOS/MS-DOS
PatentIn Release #1.0,
 Score
Pred.
                                                                                                                                                                             Sequence 2, Application US/08477531B
Patent No. 5821123
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 08/107,669 FILING DATE: 13-AUG-1993 LIOR APPLICATION DATA: APPLICATION NUMBER: PCT/US92/1
                                                                                                                                                                                    LITLE OF INVENTION: Modified A.
NUMBER OF SEQUENCES: 67
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne "
STREET: 1100
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                                                                                                                                                                                                                                                              SEE: Sterne, Kessler, 1100 New York Ave., Washington
                                                                                                                                                                                                                                                                                                STATE: D.C.
COUNTRY: United States of
ZIP: 20005-3934
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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larity 91.2%;
Conservative
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ATTORNEY/AGENT INFORMATION:
NAME: Michele A. Cimbala
REGISTPANTION:
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il Similarity 91.2%;
103; Conservative
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acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 436
PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER:
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Patent No. 6147203
GENERAL INFORMATION:
APPLICANT: Pastan, Ira
APPLICANT: Lee, Byungkook
APPLICANT: Jung, Sun-Hee
APPLICANT: Brinkmann, Ulrich
TITLE OF INVENTION: Recombinant Disulfide-Stabilized
TITLE OF INVENTION: Polypeptide Fragments Having Binding
NUMBER OF SEQUENCES: 23
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/082,842A
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                                                                                                                                                    Goldstein and Fox N.W., Suite 600
                                                                                                    Variable
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6.3e-42;
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R: 0610.1000002
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/10906
FILING DATE: 14-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/808,464
FILING DATE: 13-DEC-1991
ATTORNEY/AGENT INFORMATION:
NAME: Cimbala, Michele A.
REGISTRATION NUMBER: 33,851
REFERENCE/DOCKET NUMBER: 0610.100000;
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202/371-2600
TELEFAX: 202/371-2540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 536;
Pred. No.
                             Sequence 2, Application US/08082842A
Patent No. 5869619
GENERAL INFORMATION:
APPLICANT: Studnicka, Gary M.
TITLE OF INVENTION: Modified Antibody
NUMBER OF SEQUENCES: 89
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne, Kessler, Goldstei
                                                                                                                                                                                                                                                                                                                                                        JMBER: US/08/082,842A
23-JUN-1993
                                                                                                                                                                                                 STATE:
COUNTRY: USA
ZIP: 20005-3934
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
TYPE: IBM PC compatible
TYPE: IBM PC compatible
TYPE: IBM PC compatible
TYPE: IBM PC compatible
TYPE: IBM PC compatible
TYPE: IBM PC compatible
TYPE: IBM PC compatible
TYPE: IBM PC compatible
TYPE: IBM PC compatible
TYPE: IBM PC compatible
                                                                                                                             SSEE: Sterne, Kessler, (
T: 1100 New York Ave., N
Washington
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91.2%;
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LENGTH: 113 amino acid
TYPE: amino acid
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nes 103; Conser
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                  -842A-2
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US-09-002-753A-4
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CITY: W
STATE:
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RESULT 12
US-08-082-
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Matches
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/note= "Complementarity Determining
Region 2 (CDR2)"
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chain variable region
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           l and Crew LL
Eighth Floor
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/note= "Framework Region
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"Complementarity
(CDR1)"
                                                                                                                                                                             Version
                                                                                                                                                                                                                                                                                                                                                                015280-152100US
                                                                                              ZIP: 94111-3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Vers
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/002,753A
FILING DATE: 05-JAN-1998
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/077,252
FILING DATE: 14-JUN-1993
ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen Lauver
REGISTRATION NUMBER: 015280-1521(
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
TELEFAX: (415) 576-0300
TELEFAX: (415) 576-0300
TELEFAX: (415) 376-0300
TELEFAX: (415) 376-0300
TELEFAX: (415) 376-0300
TELEFAX: (13) amino acid
STRANDEDNESS:
TOPOLOGY: linear
MOLECULE TYPE: protein
FEATURE:
NAME/KEY: Protein
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/note= "Framework
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               Townsend
Center, E
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/note= "C
Region 1
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/note= "
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Light c
  ADDRESSEE: Townsend and I
STREET: Two Embarcadero C
CITY: San Francisco
STATE: California
COUNTRY: USA
ZIP: 94111-3834
CORRESPONDENCE ADDRESS:
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LOCATION: 1..113
OTHER INFORMATION:
OTHER INFORMATION:
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LOCATION: 1..23
OTHER INFORMATION:
OTHER INFORMATION:
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LOCATION: 24..40
OTHER INFORMATION:
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LOCATION: 41..55
OTHER INFORMATION:
OTHER INFORMATION:
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LOCATION: 56..62
OTHER INFORMATION:
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LOCATION: 63..94
OTHER INFORMATION:
OTHER INFORMATION:
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NAME/KEY: Region
LOCATION: 95..103
OTHER INFORMATION:
OTHER INFORMATION:
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NAME/KEY:
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Fragments Having Binding Specificity
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ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                  Length
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Eighth Floor
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APPLICATION NUMBER: 09/002,753
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen Lauver
REGISTRATION NUMBER: 32,762
REFERENCE/DOCKET NUMBER: 0152100US
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                           Region
                                                                  2;
-42
                                                                  536; DB
No. 6.3e-
                                                                                            Mismatches
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend
STREET: Two Embarcadero Center, E
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/09/657, 2
FILING DATE: 07-Sep-2000
CLASSIFICATION: <Unknown>
             FR4
"Framework
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                                                                  Score
Pred.
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US-09-657-274-4

Sequence 4, Application US/09657274

Patent No. 6558672

GENERAL INFORMATION:

APPLICANT: Pastan, Ira

Lee, Byungkook

Jung, Sun-Hee

Brinkmann, Ulrich

Brinkmann, Ulrich
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 113 amino acids
TYPE: amino acid
STRANDEDNESS: <Unknown>
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             /label=
/note= "
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ilarity 91.2%;
Conservative
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LOCATION: 1..23
OTHER INFORMATION:
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protein
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LOCATION: 1.113
OTHER INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERIST
                                                                                                                                                                                                                                                                                                                                                               OF INVENTION:
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LOCATION: 104..109
OTHER INFORMATION:
OTHER INFORMATION:
                                                                              Similarity
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                                       -002-753A-4
                                                                                            103;
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                                                                  / Match
Local S
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                                                                                            Matches
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Best 1
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Fragments Having Binding Specificity
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6.3e-42;
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                  nrORMATION: /label= CDR1
"Complementarity Determining
1 (CDR1)"
                                                                                                                                                                       "COMPLEMENTATION: /label= CDR2
"Complementarity Determining
2 (CDR2)"
                                                                                                                                                                                                                                                                                                 NAME/KEY: Region
LOCATION: 95..103
OTHER INFORMATION: /label= CDR3
/note= "Complementarity Determining
Region 3 (CDR3)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 536; DB
Pred. No. 6.3e
3; Mismatches
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(FR4)
                                                                                                            FR2 (FR2)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen L.
REGISTRATION NUMBER: 32,762
REFERENCE/DOCKET NUMBER: 15280-152
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 543-9600
TELEFAX: (415) 543-5043
                                                                                                                                                                                                                                                                                                                                                                                                                     /label=
                                                                                                                                                                                                                                                                /label=
                                                                                                                                                                                                                                                                                                                                                                                                                                k Region 4
SEQ ID NO:
                                                                                                                                                                                                                                    NAME/KEY: Region
LOCATION: 63..94
OTHER INFORMATION: /label:
/note= "Framework Region 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MBER: PCT/US94/066
14-JUN-1994
                                                                                                            /label
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 4, Application PC/TUS9406687
GENERAL INFORMATION:
APPLICANT:
TITLE OF INVENTION: Recombinant D
TITLE OF INVENTION: Polypeptide F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IBM PC compatible
                                                                                                                          "Framework Region
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Polypeptide
14
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                                                                          LOCATION: 41..55
OTHER INFORMATION:
/note= "Frameword"
                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: Region
LOCATION: 104..109
OTHER INFORMATION:
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Local Similarity 91.2%;
nes 103; Conservative
                                                                                                                                                                LOCATION: 56..62
OTHER INFORMATION:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                     /note= "Framework
SEQUENCE DESCRIPTION: S
                                                                                                                                                       Region
56..62
NAME/KEY: Region
LOCATION: 24..40
OTHER INFORMATION
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OPERATING SYSTEM: PC-D
SOFTWARE: PatentIn Rel
CURRENT APPLICATION DATA:
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FILING DATE: 14-JU
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Region 1
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                                                                                                                                                                    Determining Region"
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                                                                                                                                                                                                                                                                                                                     /label= CDR2
/note= "Complementarity Determining Region
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No. 6.3e-42;
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"Framework Region
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/note= "Framework Region
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/note= "Framework Region
                                                                                                                                                                                                                                                                                                                                                                                  Region
                                                                                                                                                         CDR1
                                                                                                                                                                                                                                                                                                                                                                                                                           /label= CDR3
/note= "Complementarity
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to a Cys for possible
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"Framework
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site."
 4.
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ilarity 91.2%;
Conservative
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 113 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
NAME/KEY: Modified-site
LOCATION: 106
OTHER INFORMATION: to a C
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Modified-site
93
                                                                                                                                                                                                                                           Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: Region
LOCATION: 104..113
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                      CATION: 63..94
OTHER INFORMATION:
OTHER INFORMATION:
FEATURE:
NAME/KEY: Region
LOCATION: 95
                                                                                                                                                                                                                    PEATURE:
NAME/KEY: MOGIL-
LOCATION: 49
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                 LOCATION: 95.103
OTHER INFORMATION:
FEATURE:
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LOCATION: 41..55
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                 NAME/KEY: Region
LOCATION: 1..23
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                    NAME/KEY: Region
LOCATION: 24..40
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                               NAME/KEY: Region
LOCATION: 56..62
OTHER INFORMATION:
OTHER INFORMATION:
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OTHER INFORMATION:
CT-US94-06687-4
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Best Local Similarity
Matches 103, Conser
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US-10-160-505-71

US-11-218-813-71

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US-10-239-656-59

US-10-239-656-59

US-10-743-697-1

US-10-239-656-71

US-10-239-656-79

US-10-239-656-79

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US-09-995-529-2
; Sequence 2, Application US/09995529
; Publication No. US20040091482A9
; GENERAL INFORMATION:
; APPLICANT: Watkins, Jeffry D.
; APPLICANT: Huse, William D.
; APPLICANT: Tang, Ying
; TITLE OF INVENTION: Humanized Collagen An; TITLE OF INVENTION: Related Methods
; FILE REFERENCE: P-IX 4976
; CURRENT APPLICATION NUMBER: US/09/995,529
; CURRENT FILING DATE: 2001-11-26
; NUMBER OF SEQ ID NOS: 358
; SOFTWARE: FastSEQ for Windows Version 4.0
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US-09-995-529-2
; Sequence 2, Application US/09995529
; Publication No. US20030099655A1
; GENERAL INFORMATION:
; APPLICANT: Watkins, Jeffry D.
; APPLICANT: Huse, William D.
; APPLICANT: Tang, Ying
; TITLE OF INVENTION: Humanized Collagen A:
; TITLE OF INVENTION: Related Methods
; FILE REFERENCE: P-IX 4976
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CLASSIFICATION:

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/470110

FILING DATE: 06-JUN-1995

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/467420

FILING DATE: 06-JUN-1995

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/363131

FILING DATE: 23-DEC-1994

ATTORNEY/AGENT INFORMATION: INFORMATION:

NAME: Sutton, Jeffrey A.

REGISTRATION NUMBER: 34,028

REFERENCE/DOCKET NUMBER: P50282-2
                                                                                                                Mismatches
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APPLICANT: Ames, Robert S.
APPLICANT: Appelbaum, Edward R.
APPLICANT: Chaiken, Irwin M.
APPLICANT: Cook, Richard M.
APPLICANT: Gross, Mitchell S.
APPLICANT: Holmes, Stephen D.
APPLICANT: Holmes, Stephen D.
APPLICANT: Theisen, Timothy W.
TITLE OF INVENTION: Recombinant IL5 Ant
TITLE OF INVENTION: Treatment of IL5 MC
NUMBER OF SEQUENCES: 76
CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corp
STREET: P.O. Box 1539-UW2220
CITY: King of Prussia
COUNTRY: USA
ZIP: 19406-0939
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPPRATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN NUMBER: US/10/144,644
FILING DATE: 2002-08-08
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Sequence 45, Application US/10144644
Publication No. US20030059429A1
GENERAL INFORMATION:
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TELEPHONE: 610-270-5024
TELEFAX: 610-270-5090
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 Length 113
                               Indels
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                    . 6
ore 541; DB 4;
ed. No. 6.6e-41;
Mismatches 6
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REGISTRATION NUMBER: 34,028
REFERENCE/DOCKET NUMBER: P50282-2
ECOMMUNICATION INFORMATION:
TELEPHONE: 610-270-5024
TELEFAX: 610-270-5090
ION FOR SEQ ID NO: 45:
UENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corp
STREET: P.O. Box 1539-UW2220
CITY: King of Prussia
STATE: Pennsylvania
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/10/144,644
FILING DATE: 2002-08-08
APPLICATION NUMBER: US 08/470110
FILING DATE: 06-JUN-1995
APPLICATION NUMBER: US 08/467420
FILING DATE: 06-JUN-1995
APPLICATION NUMBER: US 08/363131
FILING DATE: 23-DEC-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/700,74
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Of ILS
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                                                                                                                                                                                                                         Sequence 45, Application US/10700740; Publication No. US20040156850A1; GENERAL INFORMATION:
APPLICANT: Ames, Robert S.
Appelbaum, Edward R.
Chaiken, Irwin M.
Cook, Richard M.
Gross, Mitchell S.
Holmes, Stephen D.
McMillan, Lynette J.
Theisen, Timothy W.
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 Score
Pred.
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ZIP: 19406-0939
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEC
89.3%;
ilarity 91.2%;
Conservative
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CLASSIFICATION:
R APPLICATION DATA
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                 Local Similarity
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103; Co
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 Query Match
Best Local S
Matches 103
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Holmes, Stephen D.
McMillan, Lynette J.
Theisen, Timothy W.
INVENTION: Recombinant ILS Antagonists Useful Treatment of ILS Mediated Disorders
                                                   DIVMTQSPSSLSVSAGEKVTMSCKSSQSLLNSRNQKNYLAWYQQKPGQP
  Length 113
                            Indels
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Score 541; DB 4;
Pred. No. 6.6e-41;
; Mismatches 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/10/700,740
FILING DATE: 04-Nov-2003
APPLICATION NUMBER: US/10/144,644
FILING DATE: 2002-08-08
APPLICATION NUMBER: US 08/470110
FILING DATE: 06-JUN-1995
APPLICATION NUMBER: US 08/467420
FILING DATE: 06-JUN-1995
APPLICATION NUMBER: US 08/363131
FILING DATE: 23-DEC-1994
ATTORNEY/AGENT INFORMATION:
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NUMBER: P50282-2
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/11/076,39
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ADDRESSEE: SmithKline Beecham
STREET: P.O. Box 1539-UW2220
CITY: King of Prussia
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                                                                                                                                                                                                                                               Appelbaum, Edward F
Chaiken, Irwin M.
                                                                                                                                                                                        Sequence 45, Application US/11076395
Sequence 45, Application US/11076395
Publication No. US20060029594A1
GENERAL INFORMATION:
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TELEFAX: 610-270-5090
INFORMATION FOR SEQ ID NO: 45:
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                            4.
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TOPOLOGY: linear
CULE TYPE: protein
JENCE DESCRIPTION: SEQ
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MEDIUM TYPE: Floppy
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LENGTH: 113 amino a
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Similarity 91.2%; 33; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE: Pennsylva
COUNTRY: USA
ZIP: 19406-0939
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CLASSIFICATION:
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Query Match
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                           Gaps
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1.9e-40;
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Score 541; DB 6;
Pred. No. 6.6e-41
; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Very
RENT APPLICATION DATA:
APPLICATION NUMBER: US/10/340,189
FILING DATE: 10-Jan-2003
CLASSIFICATION DATA:
APPLICATION NUMBER: US/09/245,202A
FILING DATE: 23-JUN-1993
APPLICATION NUMBER: PCT/US92/10906
FILING DATE: 14-DEC-1992
APPLICATION NUMBER: US 07/808,464
FILING DATE: 13-DEC-1991
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TION:
Janet M.
PR: 32,918
                                                                                                                                                                                                                                          APPLICANT: Studnicka, Gary M.
TITLE OF INVENTION: Modified Antibody Va
NUMBER OF SEQUENCES: 89
CORRESPONDENCE ADDRESS:
ADDRESSEE: McAndrews, Held & Malloy
STREET: 500 W. Madison Street, 34th
CITY: Chicago
STATE: Illinois
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 536; DB
Pred. No. 1.9e
3; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: McNicholas, Janet M.
REGISTRATION NUMBER: 32,918
REFERENCE/DOCKET NUMBER: 11
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312/707-8889
TELEFAX: 312/707-9155
                                                                                                                                                                                         US-10-340-189-2; Sequence 2, Application US/10340189; Publication No. US20030229207A1; GENERAL INFORMATION:
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89.3%;
ilarity 91.2%;
Conservative
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Local Similarity 91.2%;
les 103; Conservative
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SEQUENCE DESCRIPTION: 8
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Matches
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ESGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCQNDHSYPLTFGAGTKLEIK
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FNTION: Modified Antibody Variable Domains
QUENCES: 67
                                                                                                                                                                                                                                                                                                                      #1
                                                                                                                               Variable
                                                                                                                                                                                                                                                                                                                      Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             7;
                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: McAndrews, Held & Malloy, I
STREET: 500 West Madison Street, 34th
CITY: Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/09/097,980
FILING DATE: 16-JUN-1998
APPLICATION NUMBER: 08/107,669
FILING DATE: 13-AUG-1993
APPLICATION NUMBER: PCT/US92/10906
FILING DATE: 14-DEC-1992
APPLICATION NUMBER: US 07/808,464
FILING DATE: 13-DEC-1991
                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/325,696
FILING DATE: 18-Dec-2002
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            d. No. 1.90 Mismatches
                                                                                                                                                                                                                                   America
                                                                                                               APPLICANT: Studnicka, Gary M.
TITLE OF INVENTION: Modified Antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ph.D
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Pred. No. 1
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REGISTRATION NUMBER: 32,918
REFERENCE/DOCKET NUMBER: 11(
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5: 312/707-8889
312/707-9050
                                                                    Sequence 2, Application US/10325696
Publication No. US20040005630A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                   States
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Publication No. US20050239141A1
GENERAL INFORMATION:
APPLICANT: Studnicka, Gary M.
TITLE OF INVENTION: Modified A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
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91.2%;
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CHARACTERISTIC
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nes 103; Conser
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STATE:
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                                                       US-10-325-696-2
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                                                                                                                                                                                                                         Version #1.25
             SEE: McAndrews, Held & Malloy, Ltd.
: 500 West Madison Street, 34th Floor
Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 536; DB 6;
Pred. No. 1.9e-40
3; Mismatches
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APPLICANT: ImmunoGen, Inc.
TITLE OF INVENTION: LEUKEMIA USING THE SAME; FILE REFERENCE: A8427; CURRENT APPLICATION NUMBER: US/10/700,632; CURRENT FILING DATE: 2003-11-05; PRIOR APPLICATION NUMBER: US 60/424,332; PRIOR FILING DATE: 2002-11-07; NUMBER OF SEQ ID NOS: 94; SOFTWARE: Patentin version 3.2
                                                                                                                                                                                                                                                                                              CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/097,980
FILING DATE: 16-JUN-1998
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/107,669
FILING DATE: 13-AUG-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/10906
FILING DATE: 14-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/808,464
FILING DATE: 13-DEC-1991
            ADDRESSEE: McAndrews, Held & Malloy STREET: 500 West Madison Street, 34 CITY: Chicago STATE: IL COUNTRY: United States of America ZIP: 60661
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0, Ve CURRENT APPLICATION DATA: APPLICATION NUMBER: US/11/133,775 FILING DATE: 19-MAY-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 07/808,464
FILING DATE: 13-DEC-1991
ATTORNEY/AGENT INFORMATION:
NAME: Janet M. McNicholas, Ph.D.
REGISTRATION NUMBER: 32,918
                                                                                                   America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: 1102
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312/707-8889
TELEFAX: 312/707-9050
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 113 amino acids
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illarity 91.2%;
Conservative
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CORRESPONDENCE ADDRESS
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APPLICANT: Appelbaum, Edward R.
APPLICANT: Appelbaum, Edward R.
APPLICANT: Cook, Katchard M.
APPLICANT: Cook, Katchard M.
APPLICANT: Gords, Mitchell S.
APPLICANT: Holmes, Stephen D.
APPLICANT: Holmes, Stephen D.
APPLICANT: Holmes, Stephen D.
APPLICANT: Holmes, Stephen D.
APPLICANT: Thelasen, Timchy W.
TITLE OF INVENTION: Recombinant LILS Antagonists Useful in TITLE OF INVENTION: Treatment of ILS Mediated Disorders NUMBERS OF SEQUENCES: 76
CORRESPONDENCE ADDRESS:
ADDRESSES: Smithkline Beecham Corp STREET: P.O. Box 1539-UW2220
CITY: King of Prussia
STREET: P.O. Box 1539-UW2220
CITY: King of Prussia
STREET: Both OF Compatible
COUNTRY: USA
COUNTRY: USA
COUNTRY: STREET: PLOPPY disk
COUNTRY: 1946-0939
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US 08/470110
FILING DATE: 06-UNN-1995
FILING DATE: 06-UNN-1995
FILING DATE: 06-UNN-1995
FILING DATE: 06-UNN-1995
FILING DATE: 06-UNN-1995
FILING DATE: 06-UNN-1995
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FILING DATE: 06-UNN-1995
FILING DATE: 06-UNN-1995
FILING DATE: 06-UNN-1995
FILING DATE: 06-UNN-19
                                                                                                                                               Length 115;
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                                                                                                                                        Score 536; DB 5; 1
Pred. No. 1.9e-40;
3; Mismatches 7;
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Publication No. US20030059429A1
GENERAL INFORMATION:
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                                                                                                                                            Match 88.4%;
Local Similarity 91.2%;
nes 103; Conservative
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LENGTH: 113 amino acida
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: protein
                                                                   Mus musculus
-68
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US-10-144-644-46
; LENGTH: 11
; TYPE: PRT
; ORGANISM: 1
US-10-700-632-
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Matches
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                             Indels
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
  Length
                  . 6
Score 535; DB 4;
Pred. No. 2.3e-40
; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                        Corp
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APPLICATION NUMBER: US 08/470110
FILING DATE: 06-JUN-1995

APPLICATION NUMBER: US 08/467420
FILING DATE: 06-JUN-1995

APPLICATION NUMBER: US 08/363131

FILING DATE: 23-DEC-1994

ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/10/700,740
FILING DATE: 04-Nov-2003
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US/10/144,644
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ADDRESSEE: SmithKline Beecham
STREET: P.O. Box 1539-UW2220
CITY: King of Prussia
                                                                                                                                                                                                                                                                                                                                                                                            of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Sutton, Jeffrey A.
REGISTRATION NUMBER: 34,028
REFERENCE/DOCKET NUMBER: P5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID NO:
                                                                                                                                                                                          RESULT 11
US-10-700-740-46

J Sequence 46, Application US/10700740

Publication No. US20040156850A1

GENERAL INFORMATION:

APPLICANT: Ames, Robert S.

Appelbaum, Edward R.

Chaiken, Irwin M.

Cook, Richard M.

Gross, Mitchell S.
                                                                                                                                                                                                                                                                                                                                                                               Recombinant
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TELEPHONE: 610-270-5024
TELEFAX: 610-270-5090
INFORMATION FOR SEQ ID NO: 46:
                                                                                                                                                                                                                                                                                                                      Gross, Mitchell S.
Holmes, Stephen D.
McMillan, Lynette,
Theisen, Timothy W
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pennsylvania
ch 88.3% 1 Similarity 90.3% 102; Conservative
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TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              -0939
                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCES:
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COUNTRY:
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ILS A
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Publication No. US20030059429A1

GENERAL INFORMATION:
APPLICANT: Amee, Robert S.
APPLICANT: Chaiken, Irwin M.
APPLICANT: Chaiken, Irwin M.
APPLICANT: Cook, Richard M.
APPLICANT: Cook, Richard M.
APPLICANT: Holmes, Stephen D.
APPLICANT: Holmes, Stephen D.
APPLICANT: Theisen, Timothy W.
TITLE OF INVENTION: Recombinant IL5 P
TITLE OF INVENTION: Treatment of IL5
NUMBER OF SEQUENCES: 76
CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corp
STREET: P.O. Box 1539-UW2220
CITY: King of Prussia
STATE: Pennsylvania
COUNTRY: USA
ZIP: 19406-0939
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN RELEASE
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/144,644
FILING DATE: 2002-08-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Corp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Sutton, Jeffrey A.
REGISTRATION NUMBER: 34,028
REFERENCE/DOCKET NUMBER: P50282-2
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-270-5024
TELEFAX: 610-270-5090
INFORMATION FOR SEQ ID NO: 16:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score
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              ch 88.3%;
1 Similarity 90.3%;
102; Conservative
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LENGTH: 113 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
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                  Match
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                Length 11
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ZIP: 19406-0939

COUNTRY: USA

ZIP: 19406-0939

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PATENTION DATA:

APPLICATION NUMBER: US/11/076,395

FILING DATE: 09-Mar-2005

CLASSIFICATION SATA:

APPLICATION NUMBER: US/10/144,644

FILING DATE: 04-Nov-2003

APPLICATION NUMBER: US/8/10/10

FILING DATE: 06-JUN-1995

APPLICATION NUMBER: US 08/467420

FILING DATE: 06-JUN-1995

APPLICATION NUMBER: US 08/467420

FILING DATE: 06-JUN-1995

APPLICATION NUMBER: US 08/363131

FILING DATE: 23-DEC-1994

ATTORNEY/AGENT INFORMATION:

NAME STATES OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL OF TOTAL
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              Score 535; DB 4;
Pred. No. 2.3e-40;
; Mismatches (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: SmithKline Beecham Corp
STREET: P.O. Box 1539-UW2220
CITY: King of Prussia
STATE: Pennsylvania
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       34,028
3ER: P50282-
                                                                                                                                                                                                                                                                                                                                                                                                                                                           NT: Ames, Robert S.
Appelbaum, Edward R.
Chaiken, Irwin M.
Cook, Richard M.
Gross, Mitchell S.
Holmes, Stephen D.
McMillan, Lynette J.
Theisen, Timothy W.
FINVENTION: Recombinant I
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TELEPHONE: 610-270-5024
TELEFAX: 610-270-5090
MATION FOR SEQ ID NO: 46:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Treatment: 76
                                                                                                                                                                                                                                                                                                                                                             Sequence 46, Application US/11076395
Sequence 46, Application US/11076395
Publication No. US20060029594A1
GENERAL INFORMATION:
APPLICANT: Ames, Robert S.
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REGISTRATION NUMBER: 34
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TOPOLOGY: linear
CULE TYPE: protein
ENCE DESCRIPTION: S
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CORRESPONDENCE ADDRE:
ADDRESSEE: Smit)
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Score 535; DB 6;
Pred. No. 2.3e-40;
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ed. No. 1.5e
Mismatches
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/470110
FILING DATE: 06-JUN-1995
PRIOR APPLICATION NUMBER: US 08/467420
FILING DATE: 06-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/363131
FILING DATE: 23-DEC-1994
ATTORNEY/AGENT INFORMATION: INFORMATION:
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Treatment of IL5 Mediated Disorder:
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Pred. No. 1.5e-39
; Mismatches
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CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corp
STEET: P.O. Box 1539-UW220
CITY: King of Prussia
CITY: King of Prussia
CITY: USA
ZIP: 19406-0939
COMPUTER: READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Vers
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/11/076,395
FILING DATE: 09-Mar-2005
CLASSIFICATION NUMBER: US/10/144,644
FILING DATE: 2002-08-08
APPLICATION NUMBER: US/0/144,644
FILING DATE: 06-JUN-1995
APPLICATION NUMBER: US 08/470110
FILING DATE: 06-JUN-1995
APPLICATION NUMBER: US 08/47011
FILING DATE: 06-JUN-1995
APPLICATION NUMBER: US 08/47010
FILING DATE: 06-JUN-1995
APPLICATION NUMBER: US 08/467420
FILING DATE: 06-JUN-1995
APPLICATION NUMBER: US 08/467420
FILING DATE: 06-JUN-1995
APPLICATION NUMBER: US 08/363131
FILING DATE: 06-JUN-1995
APPLICATION NUMBER: US 08/363131
FILING DATE: 06-JUN-1995
APPLICATION NUMBER: US 08/467420
FILING DATE: 06-JUN-1995
APPLICATION NUMBER: US 08/263131
FILING DATE: 06-JUN-1995
APPLICATION NUMBER: US 08/263131
FILING DATE: 06-JUN-1995
APPLICATION NUMBER: US 08/263131
FILING DATE: 06-JUN-1995
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FILING DATE: 06-JUN-1995
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APPLICATION NUMBER: US 08/263131
FILING DATE: 06-JUN-1995
APPLICATION NUMBER: US 08/263131
FILING DATE: 06-JUN-1995
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Chaiken, Irwin M.
Cook, Richard M.
Gross, Mitchell S.
Holmes, Stephen D.
McMillan, Lynette J
Theisen, Timothy W.
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Publication No. US20060029594A1
GENERAL INFORMATION:
APPLICANT: Ames, Robert S.
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INFORMATION FOR SEQ ID NO: 16:
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TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
CULE TYPE: protein
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SEQUENCE
US-11-076-395-16
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US-11-076-395-16
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Pred. No. 1.5e-3;
; Mismatches
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CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/10/144,644

FILING DATE: 2002-08-08

APPLICATION NUMBER: US 08/467420

FILING DATE: 06-JUN-1995

APPLICATION NUMBER: US 08/467420

FILING DATE: 06-JUN-1995

APPLICATION NUMBER: US 08/363131

FILING DATE: 23-DEC-1994

ATTORNEY/AGENT INFORMATION:

REGISTRATION NUMBER: 34,028

REGISTRATION NUMBER: 34,028

REFERENCE/DOCKET NUMBER: P50282-2

TELECOMMUNICATION INFORMATION:

TELEPHONE: 610-270-5024

TELEFAX: 610-270-5020
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Corp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/700,74
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CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham
STREET: P.O. Box 1539-UW2220
CITY: King of Prussia
STATE: Pennsylvania
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Appelbaum, Edward F
Chaiken, Irwin M.
Cook, Richard M.
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US-10-700-740-16
Sequence 16, Application US/10700740
Publication No. US20040156850A1
GENERAL INFORMATION:
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Holmes, Stephen
McMillan, Lynett
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SEQUENCE CHARACTERISTICS
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SEQUENCE DESCRIPTION: 3
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CITY: King of Pr
STATE: Pennsylva
COUNTRY: USA
ZIP: 19406-0939
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US-11-075-891-26
; Sequence 26, Application US/11075891
; Publication No. US20060088521A1
; GENERAL INFORMATION:
; APPLICANT: MAHADEVAN, DARUKA
; TITLE OF INVENTION: COMPOSITION AND ME;
; FILE REFERENCE: 263922US96
; CURRENT APPLICATION NUMBER: US/11/075,
; CURRENT FILING DATE: 2005-03-10
; PRIOR APPLICATION NUMBER: US 60/557,25
; PRIOR FILING DATE: 2004-03-27
; NUMBER OF SEQ ID NOS: 36
; SOFTWARE: PatentIn version 3.3 Peptide 3; Sequence U uery Match est Local Similarity 91.2%; atches 103; Conservative Syntheti SEQ ID NO 26 LENGTH: 118 TYPE: PRT ORGANISM: Artificial S FEATURE: OTHER INFORMATION: 1-075-891-26 382 382 382 382 382 382 383 383 393 373 373 373 64 マ Query Ma Best Loc Matches a 엄 8 ਠੋ  $\supset$ 

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US-11-219-563-71
; Sequence 71, Application US/11219563
; Publication No. US20060088539A1
; GENERAL INFORMATION:
; APPLICANT: Bander, Neil
; TITLE OF INVENTION: MEMBRANE ANTIGEN AND USES THEREOF
; FILE REFERENCE: 13651.001 (BZL-001)
; CURRENT APPLICATION NUMBER: US/11/219,563
; CURRENT FILING DATE: 2005-09-02
; PRIOR APPLICATION NUMBER: PCT/US04/06586
; PRIOR FILING DATE: 2004-03-03

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TYPE: PRT
ORGANISM: Artificial
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US-11-219-563-91

Sequence 91, Application US/11219563

Fublication No. US20060088539A1

GENERAL INFORMATION:

APPLICANT: Bander, Neil

TITLE OF INVENTION: MEMBRANE ANTIGEN AND USES THEREOF

TITLE OF INVENTION: MEMBRANE ANTIGEN AND USES THEREOF

TITLE OF INVENTION: MEMBRANE ANTIGEN AND USES THEREOF

FILE REFERENCE: 13651.001 (BZL-001)

CURRENT APPLICATION NUMBER: US/11/219,563

CURRENT APPLICATION NUMBER: PCT/US04/06586

PRIOR FILING DATE: 2004-03-03

PRIOR FILING DATE: 2003-03-03

PRIOR FILING DATE: 2003-03-03

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PRIOR FILING DATE: 2003-05-30

PRIOR FILING DATE: 2003-05-30
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US-11-219-563-72
; Sequence 72, Application US/11219563
; Publication No. US20060088539A1
; GENERAL INFORMATION:
; APPLICANT: Bander, Neil
; TITLE OF INVENTION: MODIFIED ANTIBODIES TO PROSTATE-SPECIFIC
; TITLE OF INVENTION: MEMBRANE ANTIGEN AND USES THEREOF
; FILE REFERENCE: 13651.001 (BZL-001)
; CURRENT APPLICATION NUMBER: US/11/219,563
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SOFTWARE: FastSEQ for Windows Version
PRIOR APPLICATION NUMBER: US 10/379,8:
PRIOR FILING DATE: 2003-03-03
PRIOR APPLICATION NUMBER: 10/449,379
PRIOR FILING DATE: 2003-05-30
NUMBER OF SEQ ID NOS: 144
SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 71
LENGTH: 113
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TITLE OF INVENTION: MODIFIED ANTIBODIES TO PROSTATE-SPECIFIC
TITLE OF INVENTION: MEMBRANE ANTIGEN AND USES THEREOF
FILE REFERENCE: 13651.001 (BZL-001)
CURRENT APPLICATION NUMBER: US/11/219,563
CURRENT FILING DATE: 2005-09-02
PRIOR FILING DATE: 2004-03-03
PRIOR FILING DATE: 2004-03-03
PRIOR FILING DATE: 2003-03-03
PRIOR FILING DATE: 2003-05-30
NUMBER OF SEQ ID NOS: 144
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6.5e-22;
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3.4e-21;
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Pred. No. 3
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                                                                                                                                                                                                                                                            Score 483;
Pred. No.
                                                                                                                        0
CURRENT FILING DATE: 2005-09-02
PRIOR APPLICATION NUMBER: PCT/US04/06586
PRIOR FILING DATE: 2004-03-03
PRIOR APPLICATION NUMBER: US 10/379,838
PRIOR FILING DATE: 2003-03
PRIOR APPLICATION NUMBER: 10/449,379
PRIOR FILING DATE: 2003-05-30
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US-10-511-937-2965
· cemuence 2965, Application US/10511937
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Sequence 92, Application US/11219563
Publication No. US20060088539A1
GENERAL INFORMATION:
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79.6%;
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                                                                                                      NUMBER OF SEQ ID NOS: 144
SOFTWARE: FastSEQ for Windows
SEQ ID NO 72
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APPLICANT: Wyeth
TITLE OF INVENTION: HUMANIZED ANTI-ST4 ANTIBODIES AND ANTI-ST4/CALICHEAMICIN CONJUGATI
FILE REFERENCE: 040000-0317285
CURRENT APPLICATION NUMBER: US/11/221,902
CURRENT FILING DATE: 2005-09-09
NUMBER OF SEQ ID NOS: 89
SOFTWARE: Patentin version 3.3
SEQ ID NO 11
LENGTH: 214
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                                        442; DB 7; L
No. 1.2e-19;
smatches 17;
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Pred. No. 5.2e-18
12; Mismatches 14
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CURRENT APPLICATION NUMBER: US/11/221,902
CURRENT FILING DATE: 2005-09-09
NUMBER OF SEQ ID NOS: 89
SOFTWARE: Patentin version 3.3
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                                             Score
Pred.
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Sequence 23, Application US/11221902
Publication No. US20060088522A1
GENERAL INFORMATION:
APPLICANT: Wyeth
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Publication No. US20060088522A1
GENERAL INFORMATION:
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sequences
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                                             th 72.9%;
Similarity 75.2%;
85; Conservative
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NAME/KEY: MISC_FEATURE
LOCATION: (1)..(107)
OTHER INFORMATION: light
  sapiens
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LENGTH: 10,

TYPE: PRT
ORGANISM: Artificial
FEATURE:
OTHER INFORMATION: ar
OTHER INFORMATION: se
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ORGANISM: Artificial
FEATURE:
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Homo
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SOFTWARE: Pat
SEQ ID NO 23
LENGTH: 107
TYPE: PRT
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; ORGANISM:
US-11-254-679-
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Best Local S
Matches 81
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APPLICANT: Wu, Herren
APPLICANT: Allan, Christian
APPLICANT: Gao, Changshou
APPLICANT: An, Ling-Ling
APPLICANT: Mao, Ling-Ling
APPLICANT: Mao, Su-Yau
APPLICANT: Mao, Su-Yau
APPLICANT: Mao, Su-Yau
APPLICANT: Mao, Su-Yau
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APPLICANT: Mao, Su-Yau
APPLICANT: Mao, Su-Yau
APPLICANT: Mao, Su-Yau
APPLICATION NUMBER: 60/650,726
PRIOR FILING DATE: 2005-03-07
PRIOR FILING DATE: 2005-03-07
PRIOR FILING DATE: 2005-03-08
PRIOR FILING DATE: 2005-03-08
PRIOR FILING DATE: 2005-03-08
PRIOR FILING DATE: 2005-09-09
PRIOR FILING DATE: 2005-09-09
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                                                  APPLICANT: Woodward, Robert
APPLICANT: Fry, Kirk
APPLICANT: Woodward, Robert
APPLICANT: Ly, Ngoc
APPLICANT: Ly, Ngoc
APPLICANT: Ly, Ngoc
APPLICANT: Ly, Ngoc
APPLICANT: Prentice, James
APPLICANT: Prentice, James
APPLICANT: Rosenberg, Steven
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR DIAGNOSING
TITLE OF INVENTION: AND MONITORING TRANSPLANT REJECTION
FILE REFERENCE: 506612000104
CURRENT APPLICATION NUMBER: US/10/511,937
CURRENT APPLICATION NUMBER: US 10/131,831
PRIOR FILING DATE: 2003-04-24
PRIOR FILING DATE: 2002-04-24
PRIOR FILING DATE: 2002-12-20
NUMBER OF SEQ ID NOS: 3117
SOFTWARE: Patentin version 3.2
SEQ ID NO 2965
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Pred. No. 3.5e-20;
D; Mismatches 15;
           GENERAL INFORMATION:
APPLICANT: EXPRESSION DIAGNOSTICS,
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; Sequence 5, Application US/1125467;
; Publication No. US20060099207A1
; GENERAL INFORMATION:
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US20060088836A1
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ilarity 77.2%;
Conservative 1
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Best Local S
Matches 88
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3-11-221-902-5
Sequence 5, Application US/11221902
Sequence 5, Application US/11221902
Publication No. US20060088522A1
GENERAL INFORMATION:
APPLICANT: Wyeth
TITLE OF INVENTION: HUMANIZED ANTI-5T4 ANTIBODIES AND ANTI-5T4/CALICHEAMICIN CONJUGATE
FILE REFERENCE: 040000-0317285
CURRENT APPLICATION NUMBER: US/11/221,902
CURRENT FILING DATE: 2005-09-09
CURRENT FILING DATE: 2005-09-09
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16;
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                                                                                                                        DB 7;
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Pred. No. 7.4e-17
                                                                                                                                                   Mismatches
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Sequence 17, Application US/11221902

Publication No. US20060088522A1

GENERAL INFORMATION:

APPLICANT: Wyeth

TITLE OF INVENTION: HUMANIZED ANTI-5T4 ANTIE

FILE REFERENCE: 040000-0317285

CURRENT APPLICATION NUMBER: US/11/221,902

CURRENT FILING DATE: 2005-09-09

NUMBER OF SEQ ID NOS: 89

SOFTWARE: Patentin version 3.3

SEQ ID NO 17

LENGTH: 107
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Pred. No.
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OTHER INFORMATION: sequence
1-221-902-17
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76.2%;
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|larity 69.0%;
|Conservative
    -10-0
                                                                                                                                      Local Similarity 76.3
nes 77; Conservative
PRIOR FILING DATE: 2002-1
NUMBER OF SEQ ID NOS: 62
SOFTWARE: Patentin Ver. 2
SEQ ID NO 57
LENGTH: 101
TYPE: PRT
ORGANISM: Homo sapiens
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ORGANISM: Artificial
FEATURE:
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Local Similarity
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ORGANISM: Artificial
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US-11-239-308-57
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ENGTH: 214
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Best Local
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Pred. No. 8.2e-18;
2; Mismatches 14;
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14;
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1.4e-17;
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; Pred. No. 1.4e-
10; Mismatches
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US-11-239-308-57
; Sequence 57, Application US/11239308
; Publication No. US2006008883A1
; GENERAL INFORMATION:
; APPLICANT: Smider, Vaughn
; APPLICANT: Larrick, James W.
; TITLE OF INVENTION: Recombinant Catalytic Po.
; FILE REFERENCE: 021216-000310US
; CURRENT APPLICATION NUMBER: US/11/239,308
; CURRENT FILING DATE: 2005-09-28
; PRIOR FILING DATE: 2003-10-09
; PRIOR FILING DATE: 2003-10-09
                                                                                                                                                                                                                                                                                                                                                                        US-11-221-902-63
; Sequence 63, Application US/11221902
; Publication No. US20060088522A1
; GENERAL INFORMATION:
; APPLICANT: Wyeth
; TITLE OF INVENTION: HUMANIZED ANTI-5T4 ANTIE
; FILE REFERENCE: 040000-0317285
; CURRENT APPLICATION NUMBER: US/11/221,902
; CURRENT FILING DATE: 2005-09-09
; NUMBER OF SEQ ID NOS: 89
; SOFTWARE: Patentin version 3.3
; SEQ ID NO 63
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Similarity 71.7%;
81; Conservative
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Best Local Similarity 76.2%;
Matches 77; Conservative
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OTHER INFORMATION: CDR3
           NAME/KEY: MISC_FEATURE
LOCATION: (24)..(34)
OTHER INFORMATION: CDR
                                                      FEATURE:
NAME/KEY: MISC_FEATURE
LOCATION: (50)..(56)
                                                                                              OTHER INFORMATION: CDR
                                                                                                                      NAME/KEY: MISC_FEATURE
LOCATION: (89)..(97)
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Sequence 9, Application US/11221902
Publication No. US20060088522A1
GENERAL INFORMATION:
APPLICANT: Wyeth
TITLE OF INVENTION: HUMANIZED ANTI-5T4 ANTIBODIES AND ANTI-5T4/CALICHEAMICIN CONJUGAT
FILE REFERENCE: 040000-0317285
CURRENT APPLICATION NUMBER: US/11/221,902
CURRENT FILING DATE: 2005-09-09
NUMBER OF SEQ ID NOS: 89
SOFTWARE: Patentin version 3.3
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13; Mismatches 16;
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Pred. No. 1.2e-16
13; Mismatches 16
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LOCATION: (1)._(107)
OTHER INFORMATION: light chain variable region
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TITLE OF INVENTION: HUMANIZED ANTI-5T4 ANTIE
FILE REFERENCE: 040000-0317285

CURRENT APPLICATION NUMBER: US/11/221,902

CURRENT FILING DATE: 2005-09-09

NUMBER OF SEQ ID NOS: 89

SOFTWARE: PatentIn version 3.3

SEQ ID NO 7

LENGTH: 214
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OTHER INFORMATION: artificial sequence;
OTHER INFORMATION: sequences
FEATURE:
NAME/KEY: MISC_FEATURE
LOCATION: (1)..(107)
OTHER INFORMATION: light chain variable
US-11-221-902-5

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nes 78; Conservative
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llarity 69.0%;
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FEATURE:
OTHER INFORMATION: artificial sequence
OTHER INFORMATION: sequences
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Similarity 69.0%;
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LOCATION: (1)..(107)
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